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ALIGNMENTS

RESULT 1 AAY08661 09-AUG-1999 AAY08661; Human Zcytol0 allele 1 protein. AAY08661 standard; protein; 176 (first entry) **A**

Zcytol0; four alpha helix cytokine; vulnerary; cytostatic; haemostatic; anti-inflammatory; anti-asthmatic; growth factor; maintenance factor; trachea; salivary gland; stomach; panoreas; muscle; gene therapy; cancer; proliferation; differentiation; modulator; gastrointestinal tract; oral cavity; asthma; tracheobronchial tract; bronchitis; wound healing; platelet count; thrombocytopenia; human. 26-NOV-1997; 25-NOV-1998; 03-JUN-1999 WO9927103-A1 Homo sapiens 97US-00979156 98WO-US025228.

WPI; 1999-357840/30. N-PSDB; AAX77692. Conklin DC, Haldeman BA, Grossmann

A;

(ZYMO) ZYMOGENETICS INC.

New mammalian cytokine-like polynucleotide useful in the treatment of asthma and bronchitis.

Claim 1; Page 64-65; 82pp; English.

This invention describes a novel mammalian four alpha helix cytokine designated Zcytol0 which has vulnerary, cytostatic, anti-inflammatory, anti-asthmatic and haemostatic activity. Zcytol0 may be a growth and/or maintenance factor in the trachea and salivary glands, stomach, pancreas and muscle. Zcytol0 polynucleotides are useful as sources of primers and probes, and to determine if the Zcytol0 gene is present on chromosome 1, or to detect any mutations that may have occurred. Zcytol0 polynucleotides are also useful as gene therapy reagents. The Zcytol0 pancreas and

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CC polypeptide, its fragments, and antibodies, as well as compounds
CC identified as having binding affinity to Zcytol0, should be useful in the
CC treatment of conditions associated with abnormal physiology or
CC development, including abnormal proliferation, e.g. cancerous conditions,
CC or degenerative conditions or altered immunity. The products are also
CC cytokine production in the prevention or treatment of conditions
CC characterized by abnormal cell proliferation, cell differentiation or
CC cytokine production. Zcytol0 polypeptides, agonists and antagonists may
CC characterized by useful in the regeneration of the gastrointestinal
CC tract or oral cavity, as well as in the treatment of asthma, and other
CC diseases of the tracheobronchial tract such as bronchitis. Zcytol0
CC polypeptides may also be useful to promote wound healing, and to increase
CC platelet count, e.g. in cancer patients who experience thrombocytopenia
CC due to chemotherapy or radiation therapy
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                                                    Isolated subunits treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interleukin 20; IL-20; IL-20RA; ZcytoR7; IL-20RB; DIRS1; immunoglobulin;
antiinflammatory; antipsoriatic; antiasthmatic; antibacterial;
dermatological; antiulcer; antagonist.
                                                                                                                                WPI; 2001-398320/42.
                                                                                                                                                                  Foster DC,
Rixon MW,
                                                                                                                                                                                                                                                         23-DEC-1999; 99US-00471774.
22-JUN-2000; 2000US-0213416P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human interleukin-20 (IL-20) receptor polypeptide.
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                    Disclosure;
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                                                        inflammatory diseases
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                                                                         IL-20RA and
                                                                                                                                                                Presnell SR,
                                                                                                                                                                                 Xu W,
                  Page 53-54; 119pp; English
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100.0%; Pred. No. 5e-88;
:ive 0; Mismatches
                                                                       20 soluble receptor comprising two polypeptide IL-20RB, useful for down-regulating IL-20 and thus
                                                                                                                                                                Fox BA;
                                                        such as psoriasis
                                                                                                                                                                                   Kelly JD,
                                                                                                                                                                                   Sprecher CA,
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                                                                                                                                                                                                                                                                                                                                                        Human; interleukin-20; II-20; antagonist; psoriasis; eczema; de adult respiratory disease; asthma; bronchitis; pneumonia; multiple organ failure; inflammatory lung injury; septic shock; bacterial pneumonia; inflammatory bowel disease; rheumatoid art ulcerative colitis; Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammatory diseases such as psoriasis, inflammatory lung injury such asthma or bronchitis, adult respiratory disease (AD), septic shock, multiple organ failure, bacterial pneumonia, eczema, atopic and contact dermatitis, and inflammatory bowel disease such as ulcerative colitis a Crohn's disease. Sequences AAB85259-262 represent human II-20 receptor
                                                                                                                   Thompson P,
Sprecher CA,
                                                                                                                                                                                            23-DEC-1999;
22-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human interleukin-20, IL-20, protein #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAU04049;
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Blumberg
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100.0%; Pred. No. 5e-88;
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Treating interleukin-20 induced inflammation in a mammal, such as adult respiratory disease, eczema, psoriasis, contact dermatitis, multiple organ failure and septic shock, involves administration or an administration or a

WPI; 2001-418045/44

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Matches 176
                                                                                                                                                                                                                                                                                             Inflammation; interleukin-20; IL-20; interleukin-8; IL-8; chemokine; neutrophil; monocyte; basophil; eosinophil; chemoattractant; psoriasis; periodontal disease; rheumatoid arthritis; idiopathic pulmonary fibrosis; angiogenesis-dependent chronic inflammatory condition; lung cancer; melanoma; inflammatory disease; diabetes; arteriosclerosis; cataract; reperfusion injury; cancer; meningitis; rheumatic disease; skin disease; idiopathic pulmonary fibrosis; inflammatory bowel disease; psoriasis; ulcerative colitis; eczema; atopic dermatitis; contact dermatitis; ulcerative colitis; eczema; atopic dermatitis; contact dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to treating a mammal afflicted with a disease which an interleukin-20 (II-20) polypeptide plays a role, involves administering antagonist of II-20 polypeptide to the individual. The method is useful for treating psoriasis, eczema, atopic dermatitis, contact dermatitis, adult respiratory disease, asthma, bronchitis are pneumonia and is also useful for treating multiple organ failure, inflammatory lung injury, septic shock, bacterial pneumonia, inflamm bowel disease, rheumatoid arthritis, ulcerative colitis and Crohn's
                                                                (FOST/)
(XUWW/)
(MADD/)
               (SPRE/)
(BLUM/)
(EAGA/)
                                                                                                                                               23-DEC-1999;
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                                                                                                                                                                                                                                                                                   bronchitis;
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                                                                                                                                    22-JUN-2000;
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XU W.

MADDEN K L.

KELLY J D.

SPRECHER C A.

BLUMBERG H.

EAGAN M A.

JASPERS S R.
                                                                                           THOMPSON
FOSTER D
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                                                                                                                                                                                                                                                                                   pneumonia.
                                                                                                                                  99US-0171969P
2000US-0213341P
                                                                                                                                                                                                                                                                                                                                                                                                                                  disease related human interleukin (IL)20 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No. 5e-88;
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Sequence 176 AA;

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CC comprising administering antagonist of the II-20 polypoptide to the CC individual. An important cytokine in the inflammatory process is CC interleukin-8 (II-8), a chemokine in that acts as an agonist for neutrophils CC via chemotaxis and the release of granule enzymes. II-8 binds to CR receptors on neutrophils, monocytes, basophils, and eosinophils. II-8 is CR a potent chemoattractant for neutrophils; and the early stages of CR periodontal disease are characterised by the influx of neutrophils. II-8 is an other inducer of angiogenesis in several angiogenesis-dependent CR chronic inflammatory conditions, including theumatoid arthritis, CR psoriasis, and idiopathic pulmonary fibrosis. Additionally, II-8 is an CR expression correlates with experimental metastatic activity of some cell lines. Therefore an effective method to treat inflammatory diseases would be to administer an agent that would inhibit II-8. It has been shown that II-20 up-regulates II-8. Therefore antagonists to II-20 can be used to treat these diseases. The method is used for treating CR diseases including diabetes, reteriosclerosis, cataracts, reperfusion ciplury, cancer, meningitis, rheumatic diseases, idiopathic pulmonary cR (psoriasis, eczema, atopic dermatitis and contact dermatitis) or an inflammatory lung disease (adult respiratory disease (ARD), asthma, CR bronchitis and pneumonia). This sequence represents a human interleukin-CR (II-20) polypoptide used in developing the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thompson
Blumberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 12-13; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating inflammatory skin and lung diseases using antibodies against interleukins (IL)-20 (which indirectly modulates activation of IL-8),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-507215/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CHAN/) CHANDRASEKHER Y (NOVA/) NOVAK J E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for treating e.g. psoriasis, asthma and bronchitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ξ, Ţ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Foster DC,
Eagan MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                n describes a method (I) which an interleukin-20
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Jaspers
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SR, Ch
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for treating a mammal afflicted with (IL-20) polypeptide plays a role
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Chandrasekher YA,
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Novak JE;
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RESULT 5
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Best Local Sim
Matches 176;
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                                                                        AAE29052;
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                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                       DIRILARTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
                                                                                                                                                                                                                   MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
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                                                     (first
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                                                     entry)
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Pred. No. 5e-88;
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Human; heterodimeric cytokine receptor; interleukin-22R; IL-22R; asthma; inflammatory disease; psoriasis; adult respiratory disease; bronchitis; septic shock; multiple organ failure; inflammatory lung injury; eczema;

Human

IL-20

protein

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RESULT 6
ABR62464
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Soluble heterodimeric cytokine receptor useful for interleukin-20 and treating inflammatory diseases, asthma, comprises an interleukin-22R subunit and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bacterial pneumonia; dermatitis; ulcerative colitis; Crohn's disease; antiinflammatory; dermatological; antibacterial; immunosuppressive;
                                                   Homo sapiens.
                                                                          Human;
                                                                                                   Human
                                                                                                                             06-NOV-2003
                                                                                                                                                    ABR62464
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21-JUN-2001;
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                                                                                                                                                                        ABR62464 standard; protein; 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ZYMO ) ZYMOGENETICS INC
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176;
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                                                                           interleukin-20;
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2001US-0299865P
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1. .24
/label= Signal_peptide
                          Location/Qualifiers
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                                                                           IL-20;
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Pred. No. 5e-88;
Mismatches 0;
                                                                           cervical
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                                                                           cancer;
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                                                                           cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                            Length 176;
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                                                                                                                                                                                                                                                                                                                                                                                       Indels
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such as psoriasis
interleukin-20RB
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                                                                            virucide
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                Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is the protein sequence of human interleukin-20 (IL-20), including the signal sequence. The invention relates to the use of IL-20 for treating cervical cancer or cells infected with human papilloma virus. IL-20 can be administered alone or in conjunction with radiation or chemotherapeutic agents or surgical excision of the involved cells or or chemotherapeutic agents or surgical excision of the involved cells or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inhibiting the growth and/or proliferation of cervical cancer cells, useful for treating cervical cancer or human papilloma virus infection, comprises bringing Interleukin-20 (IL-20) into contact with the cervica
                                                                                                  autoimmune disease; insulin-dependent diabetes mellitus; immune response; rheumatoid arthritis; multiple sclerosis; infection; radiation therapy; asthma; bronchitis; wound healing; periodontal disease; antiinflammatory; psoriasis; eczema; dry skin; oral cavity; muscle atrophy; haemostatic;
                                                                                                                                           Human; cytokine-like polypeptide-10; Zcyto10; therapy; thrombocytopaenia; autoimmune disease; insulin-dependent diabetes mellitus; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer cells.
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                                                                                     cancer;
                                                                                                                                                                                    Human cytokine-like polypeptide-10 (2cyto10)
                                                                                                                                                                                                                                              ABW00889;
                                                                                                                                                                                                                                                                         ABW00889 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein
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                                                                                      vulnerary.
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Location/Qualifiers
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/label= Signal_peptide
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Pred. No. 5e-88;
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120 60 60

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                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated human cytokine-like polypeptide-10 CC (Zcytol0) polypeptide. Zcytol0 is useful for prevention or treatment of CC conditions characterised by improper cell proliferation, cell CC differentiation or cytokine production. Zcytol0 is useful for treating CC rheumatoid arthritis, multiple sclerosis, etc., by inhibiting cancer cell immune response. The invention is useful for inhibiting cancer cell CC growth or proliferation, for stimulating immune system to combat CC microbial or viral infections, for increasing platelet production, and thus useful in cancer patients who experience thrombocytopaenia due to cancer therapy or radiation therapy. Zcytol0 is useful in tracheases specific or tracheabronchial specific applications, such as in the concert therapy or radiation therapy. Zcytol0 is useful in trachea specific or tracheabronchial specific applications, such as in the concert therapy or radiation therapy. Zcytol0 only mucous production or mucociliary cells cunderlying the same, in regulating mucous production or mucociliary cells cunderlying the same, in regulating mucous production may also enhance wound the same of the tracheabronchial tract. Zcytol0 may also enhance wound cells can be capacially useful in the treatment of affected tissues which may be capacially useful in the treatment of periodontal disease. Zcytol0 can be capacially can dor modulating muscle tone in the tracheabronchial tract, and for treating muscle atrophy in the elderly, sick or bed-ridden. Cc capacial also useful for promoting wound healing. The present sequence cis human Zcytol0 protein
                                                                                                                                                                                                              Query Match
Best Local S
Matches 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human cytokine-like polypeptide-10, Zcyto10, useful for treating insulin-dependent diabetes mellitus, rheumatoid arthritis, multiple sclerosis, asthma, psoriasis and inhibiting cancer cell growth.
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Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Col 31-32; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-799828/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conklin
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25-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAY-1999;
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 121
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KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEETE
                      KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEETE
                                                                   DIRILARTESLODTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
                                                                                           DIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
                                                                                                                                        MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
                                                                                                                                                                                                                                                                                                                         Zcyto10
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/note=
33. .12
80. .13
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.132
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100.0%; Pred. No. 5e-88;
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                                                                                                                                                                                                                                                DB 7;
                                                                                                                                                                                                                                                Length 176;
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176
                                   176
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RESULT 8

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μ

Query Match Best Local Sim Matches 176;

Similarity

100.0%;

Conservative

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Score 910; DB 7 Pred. No. 5e-88; ; Mismatches

DB 7; 0;

Length 176; Indels

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Gaps

60 60

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61

DIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK 120

MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQBIRNGFSDIRGSVQAKDGNI MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI

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ADG46669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-NOV-1997;
25-NOV-1998;
17-MAY-1999;
                                                                                                                                                  The present invention relates to novel mammalian cytokine-like polypeptide-10 (Zcyto10) proteins and polynucleotides encoding such proteins. Sequences of the invention are useful for prometing the ho of wounds and for stimulating the proliferation of platelets. In particular Zcyto10 polypeptides may be used in trachea-specific or tracheobronchial-specific applications such as maintenance or wound repair of the tracheobronchial epithelium or cells underlying it; it
                                        regulating mucous production; mucocilary clearance of debris; in treatment of asthma, bronchitis and other diseases of the tracheobronchial tract. They are also used to promote regeneration of affected tissues which may be especially useful in the treatment of periodontal disease. Zcyto10 polypeptides can be used to treat skin conditions such as psoriasis, eczema and dry skin in general. They are also used in protein therapy. The present sequence is human Zcyto10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytokine-like polypeptide-10; Zcyto10; wound healing; platelet proliferation; wound repair; mucous production; asthma bronchitis; tracheobronchial tract disease; periodontal disease skin condition; psoriasis; eczema; dry skin; protein therapy; h vulnerary; respiratory; antibronchitic; dental; dermatological.
Sequence 176 AA,
                                                                                                                                                                                                                                                                                                            Novel isolated mammalian cytokine-like polypeptide-10 (Zcyto10), useful for treating asthma, bronchitis and other tracheobronchial damage.
                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 2;
                                                                                                                                                                                                                                                                                                                                                               N-PSDB; ADG46668
                                                                                                                                                                                                                                                                                                                                                                                                            Conklin DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-APR-2003; 2003US-00413661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003176657-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                          (CONK/) CONKLIN D C. (HALD/) HALDEMAN B A.
                                                                                                                                                                                                                                                                                                                                                                              2003-863865/80.
                                                                                                                                                                                                                                                                                                                                                                                                           Haldeman
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98US-00199586.
99US-00313458.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "Human mature Zcyto10 protein"
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                                                                                                                                                                                                                                                                               31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                            BA;
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                                  n Zcytol0
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22-JUN-2000;
22-DEC-2000;
         The invention relates to a novel method of reducing or treating inflammation in a mammal which comprises administering an antibody, antibody fragment or single chain antibody which specifically binds to a receptor of interleukin (IL)-20 comprising an IL-20RA subunit and an IL-20RB subunit. The method of the invention has antiinflammatory, antiasthmatic, antiulcer, antipsoriatic, antiarthritic, respiratory, antiasthmatic, antiulcer, antibacterial, immunosuppressive and gastrointestinal applications and may be useful for reducing or treating an inflammatory, skin disease such as psoriasis, eczema, atopic including an inflammatory skin disease such as psoriasis, eczema, atopic
                                                                                                                                                                                                                                                             Thompson
Blumberg
                                                                                                                                                                    Reducing or treating inflammation, e.g. inflammatory lung disease, comprises administering an antibody, antibody fragment or single chain antibody that specifically binds to an interleukin (IL)-20RA subunit of an IL-20 receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2004005320-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ulcerative colitis; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IL-20RB subunit; antiinflammatory; dermatological; antipsoriatic;
antiarthritic; respiratory; antiasthmatic; antiulcer; antibacterial;
immunosuppressive; gastrointestinal; skin; psoriasis; eczema;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human interleukin IL-20 protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-MAY-2004
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                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                             (THOM/)
(FOST/)
(XUWW/)
(MADD/)
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(SPRE/)
                                                                                                                                                                                                                                                                                                      (NOVA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation;
                                                                                                                                                                                                                                                                                                                  (CHAN/)
                                                                                                                                                                                                                                                                                                                                (Jasp/)
                                                                                                                                                                                                                                                                                                                                          EAGA/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                               JASPERS S R.
CHANDRASEKHER
                                                                                                                                                                                                                                                                                                                                                    SPRECHER C A.
BLUMBERG H.
                                                                                                                                                                                                                                                                                                      NOVAK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
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                                                                                                                                                                                                                                                             Foster DC,
Eagan MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0171969P.
2000US-0213341P.
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                                                                                                                                             69pp; English
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                                                                                                                                                                                                                                                             Madden KL, Kelly JD,
SR, Chandrasekher YA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interleukin; IL-20; IL-20RA subunit;
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ısis, eczema,
lung disease
                                                                                                                                                                                                                                                              Sprecher CA;
Novak JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pneumonia,
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inflammatory

Query Match Best Local S Matches 176

Local Similarity nes 176; Conserv

100.0%; ilarity 100.0%; Conservative 0

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Score 910; DB 8 Pred. No. 5e-88; ; Mismatches

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Length 176;

Indels

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Gaps

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Sequence

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RESULT 10
ADM95041
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Best Local (
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This invention relates to a novel agent for prevention and/or treatment of colon cancer, comprising of the interleukin 20 (II-20) receptor alpha chain, its partial peptide or salt. The invention may be useful for the development of compounds with a cytostatic activity. The invention is useful for preventing and/or treating colon cancer. A diagnostic agent is useful for diagnosis of colon cancer. By using II-20, II-20 receptor, its salt or its peptide, compounds that have altered binding property towards them can be efficiently screened. The present sequence is that of the human Interleukin-20 protein which is related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            colon
IL-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adult respiratory distress syndrome, asthma, bronchitis and pneumonia, well as arthritis, septic shock, multiple organ failure, inflammatory bowel disease, ulcerative colitis or Crohn's disease. The current sequence is that of the human IL-20 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-320878/30.
N-PSDB; ADM95042.
                                                                                                                                                                                                                                                                                            Agent useful interleukin :
                                                                                                                                                                                                                            Disclosure; SEQ ID NO 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-AUG-2002; 2002JP-00234880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                             TAKE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            receptor alpha chain; cytostatic; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interleukin-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; interleukin 20 receptor alpha chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13
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                                                                                                                                                                                                                                                                                         l for prevention and/or treatment 20 (IL-20), IL-20 receptor alpha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
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                                                                                                                                                                                                                          50pp; Japanese.
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0; Mismatches
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                                                                                                                                                                                                                                                                                            of colon cancer, comprises chain, IL-20 receptor beta
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The present invention provides mammalian cytokine-like (Zcyto10) proteins and their encoding polynucleotides.
                                                                                                            Claim 1; SEQ ID NO 2; 31pp; English.
                                                                                                                                                                                                                     New Zcyto10 polypeptides and polynucleotides, useful for treating autoimmune diseases (e.g. insulin dependent diabetes mellitus, multiple sclerosis or rheumatoid arthritis), or skin conditions (e.g. psoriasis or conditions)
                                                                                                                                                                                                                                                                                                                                                                                N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conklin
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25-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-OCT-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ( OMYZ)
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DB; ADQ88066.
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98US-00199586.
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polypeptide-10
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N-PSDB; ADR16327 WPI; 2004-580197/56

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Best Local 9
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Disulfide-bond
Disulfide-bond
                                                                                        26-NOV-1997;
25-NOV-1998;
15-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful for treating conditions characterised by improper cell proliferation, cell differentiation or cytokine production, specifically for treating autoimmune diseases such as insulin dependent diabetes mellitus, multiple sclerosis and rheumatoid arthritis and for inhibiting cancer cell growth. The invention is also used to stimulate the immune system, to enhance wound healing and promote regeneration of affected tissues, to increase platelet count and to treat asthma and skin conditions such as psoriasis, eczema and dry skin. The invention acts as an antidiabetic, neuroprotective, antirheumatic, antiarthritic, cytostatic, immunostimulant, vulnerary, antiasthmatic, antipsoriatic and dermatological agent. The invention is also used in gene therapy. The present sequence is human Zcytol0 longer form protein.
                                      Conklin
                                                                                                                                          27-FEB-2004;
                                                                                                                                                                    05-AUG-2004.
                                                                                                                                                                                            US2004152878-A1
                                                                                                                                                                                                                                                                                                                                                                        Cytokine-like polypeptide-10; Zcyto10; therapy; asthma; infection; psoriasis; eczema; dry skin; wound healing; platelet proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                Human
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98US-00199586.
2003US-00413661.
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                                        Haldeman
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33. .126
80. .132
81. .134
                                                                                                                                                                                                                                                                     1. .24
/label= Signal_peptide
25. .176
                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                polypeptide-10 (Zcyto10) long form protein
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Pred. No. 5e-88;
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RESULT 13
ADS18363
ID 18363
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XX ADS18
AC ADS18
AC ADS18
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AC ADS18
CW ZCYTC
KW Cell
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zcytol0; cytokine-like polypeptide; cell proliferation; cell differentiation; autoimmune disease; diabetes; multiple scl rheumatoid arthritis; asthma; psociasis; cancer; wound; gene the immunosuppressive; antidiabetic; neuroprotective; antirheumatic; antiarthritic; antiasthmatic; antipsoriatic; cytostatic; vulnera
    Region
                                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                      Disulfide-bond
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121
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Pred. No. 5e-88;
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RESULT 14
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ID ADV66
XX ADV66
XX ADV66
XX ADV66
XX Humar
XX Humar
XX Derma
KW Inter
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Best Local S
Matches 176
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25-NOV-1998; 98US-00199586
15-APR-2003; 2003US-00413661
                                                                                                                               Dermatological; interleukin-20;
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HALDEMAN B A.
GROSSMANN A.
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llarity 100.0%;
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                                                                                                                            Antiinflammatory; IL-20; cytokine;
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Pred. No. 5e-88;
Mismatches
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                                                                                                                               ; Interleukin-20 antagonist; dermatological disease; inflammation
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Matches 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to alternatively spliced interleukin (II)-CC 20 variants (I; ADV66885 and ADV66902) and promoter sequences (I; ADV66887-ADV66901). IL-20 is a member of the IJ-10 family of cytokines. CC ADV66887-ADV66901). IL-20 is a member of the IJ-10 family of cytokines. CC Members in this family are important in controlling inflammation cresponses. (I) is useful for increasing the level of II-6. Tumor Necrosis Factor (TNF) alpha, Keratinocyte Growth Factor-1 (KGF-1) or reactive coxygen species in a cell, which involves contacting (I) with a cell. (II) is useful for identifying a compound for treating an II-20-induced clisease, which involves contacting a compound with a cell containing a nucleic acid comprising II-20 promoter operably linked to a reporter gene in the compound, where the expression level of the reporter gene in the presence of the compound, if lower than that in the absence of the compound, if lower than that in the absence of the compound, if lower than that in the absence of an inflammatory disease. The II-20-induced disease is a skin disease or an inflammatory chicked was used in an alignment with the alternatively spliced human II-20 variant sequence, which was used in an alignment with the alternatively spliced human II-20 variants.
                                  gastrointestinal
                                                Therapy; cns-gen.; respiratory-gen.; antiasthmatic; antipsoriatic; antiinflammatory; psoriasis; dermatological disease; immune disorder; pulmonary disease; asthma; inflammation; respiratory disease; bronchitis; antiinflammatory; inflammation; cystic fibrosis;
                                                                                                                                      Human
                                                                                                                                                                                                                                           ADW64518
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                                disease; chemotherapy; interleukin.
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                                                                                                                                                                     entry)
                                                                                                                                      seqid:1.
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Query Match
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Matches 176;
                                                                                                                                                                                                                                                           Sequence 176 AA;
                                                                                                                                                                                                                                                                                    The present invention relates to an soluble receptor having an interleukin 20 receptor alpha (IL-20RA) subunit and an IL-20RB subunit. The invention is useful for treating inflammatory diseases such as psoriasis and inflammatory lung diseases such as asthma, bronchitis and cystic fibrosis. The present sequence is the human IL-20 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated soluble receptor comprised of an interleukin 20 receptor alpha (IL-20RA) subunit and a IL-20RB subunit, useful for treating inflammatory diseases e.g., psoriasis and inflammatory lung diseases
                                                                                                                                                                                                                                                                                                                                                                                                                     e.g.,
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22-JUN-2000; 2000US-0213416P.
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(FOXB/)
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KELLY J D.

SPRECHER C A

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FOX B A.
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             KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEFQAAVVKALGELDILLQWMEETE 176
                                                                              DIRILARTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK 120
                                                                                                                                            MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVTATNLQETRNGFSDTRGSVQAKDGNT
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ilarity 100.0%;
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1 MKASSLAFSLLSAAFYLLWT.....AVVKALGELDILLQWMBETE 176
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445.636 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	œ	7	თ	v	4	ω	2	ב	No.	Result	
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hypothetical prote	hypothetical prote	late protein, 100K	membrane export pr	sensory box histid	udp-n-acetylmurama	ATP-dependent DNA	probable exonucle	hypothetical prot	probable pre-mRNA	ATP-dependent RNA	probable synechocy	hypothetical prot	probable membrane	2-isopropylmalate		hypothetical prote	m	г.		probable beta-adap	geran	CO.		-10	interleukin 10 hom	interleukin 10 -	5	interleukin-10 pre	Description		

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A;Gene: A;Cross A;Map p A;Intro C;Super	A; Scacus: particule to A; Molecule to A; Residues: A; Cross-reformation: C; Genetics:	R;Platz DNA Seq A;Title A;Refer A;Acces	A;Description A;Reference i A;Recession: A;Status: pro A;Molecule t; A;Residues: i A;Cross-refe	A; Resid A; Resid A; Cross A; Cross R; Kube, submitt	A;Title: Disu A;Reference n A;Contents: a R;Sanjanwala, submitted to A;Reference n A;Accession:	A;Refer A;Acces A;Molec A;Resic A;Cross R;Winds	C;Speci C;Date: C;Acces R;Vieir Proc. N A;Title	RESULT A38580 interle	د م م م م م ۷ د م م م م ۷ د م م
A;Gene: CDB:IL10; IL-10 A;Cross-references: GDB:128636; OMIM:124092 A;Map position: 1q31-1q32 A;Introns: 55/3; 75/3; 126/3; 148/3 C;Superfamily: interleukin-10	A; Neacus: preliminary; translated from GB/ A; Molecule type: DNA A; Residues: 1-6 <pla> A; Cross-references: UNIPARC: UPI000011E9B9; C; Genetics:</pla>	R;Platzer, C.; Volk, H.D.; Platzer, M. DNA Seq. 4, 399-401, 1994 A;Title: 5' noncoding sequence of huma A;Reference number: 137890; MUID:95143 A;Accession: 137890	A;Description: Isolation A;Reference number: S491: A;Accession: S49110 A;Status: preliminary A;Wolecule type: DNA A;Residues: 1-6,'PVAWSS' A;Cross-references: UNIP)	A; Steatus: translated from GB/EMBL/DDBJ A; Molecule type: DNA A; Residues: 1-178 < SAN> A; Residues: 1-178 < SAN> A; Cross-references: UNIPARC: UPI0000034E50; R; Kube, D.; Platzer, C.; von Knethen, A.; submitted to the EMBL Data Library, March	A;Title: Disulfide bond as A;Reference number: A48693; A;Contents: annotation; dir A;Contents: annotation; B;Sanjanwala, B. R;Sanjanwala, B. Submitted to the EMB1 Data B;Reference number: G07695 A;Reference num	A;Reference number: A38 A;Accession: A38580 A;Accession: A38580 A;Molecule type: mRNA A;Residues: 1-178 <vie> A;Cross-references: UNI A;Cross-references: UNI A;Crobemistry 32 8807-8</vie>	C;Species: Homo sapiens (man) C;Date: 14-Feb-1992 #sequence revision 14-F C;Accession: A38580; G01539; S49110; 137890 R;Vieira, P.; de Waal-Malefyt, R.; Dang, M.; Proc. Natl. Acad. Sci. U.S.A. 88, 1172-1176 A;Title: Isolation and expression of human	RESULT 1 A38580 interleukin-10 precursor -	72.5 72.5 72.7 72.7 72 72 72
IL-10 8: GDB: q31-1q3 75/3; 1	nary; c DNA PLA> s: UNIF	1k, H.I 01, 199 ding se r: 1378	olation o r: S49110 10 nary DNA PVAWSS' <:	ted fro	e bond r: A486 ation; EMBL Da	r: A389 80 RNA VIE> S: UNII Syto, F	apiens 92 #sec 80; G01 Waal-Ma Sci. U	ecurson	77.99000
128636 2 26/3; in-10	rranslaced).; Pla	of th	M GB/E	assign 593; MU disulf ta Lib	580; MU PROT: P2	(man) quence [539; S lefyt,].S.A. J.S.A.	C - human	2581 2581 255 255 279 351
5; OMI 148/3	1000	of D:9	e hu	MBL/	ment ID:9 ide	ID:9	revi 4911 R.; 88,	an h	7 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6
IM:124092 3	IIOM GB/EMBL/DDBO	R;Platzer, C.; Volk, H.D.; Platzer, M. DNA Seq. 4, 399-401, 1994 A;Title: 5' noncoding sequence of human IL-10 gene obtained by oligo-cassette A;Reference number: 137890; MUID:95143580; PMID:7841462 A;Accession: 137890	A;Description: Isolation of the human interleukin-10-promoter. Chara A;Reference number: S49110 A;Accession: S49110 A;Accession: S49110 A;Status: preliminary A;Status: preliminary A;Molecule type: DNA A;Molecule type: DNA A;Residues: 1-6, 'PVAWSS' <kub> A;Residues: 1-6, 'PVAWSS' <kub></kub></kub>	A;Status: translated from GB/EMBL/DDBJ A;Status: translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-178 <san> A;Residues: 1-178 <san> A;Cross-references: UNIPARC:UPI0000034E50; EMBL:U16720; NID:g1041812; PI R;Kube, D.; Platzer, C.; von Knethen, A.; Straub, H.; Hafner, M.; Tesch, submitted to the EMBL Data Library, March 1994</san></san>	A;Title: Disulfide bond assignments and secondary structure an A;Reference number: A48693; MUID:93372085; PMID:8364028 A;Contents: annotation; disulfide bonds in recombinant protein R;Sanjanwala, B. R;Sanjanwala, B. Submitted to the EMBL Data Library, October 1994 B;Reference number: G07695	A;Reference number: A38580; MUID:91142134; PMID:1847510 A;Accession: A38580 A;Accession: A38580 A;Accession: A38580 A;Molecule type: mRNA A;Residues: 1-178 <vie> A;Residues: 1-178 <vie> A;Residues: 1-178 <vie> A;Cross-references: UNIPROT:P22301; UNIPARC:UPI0000034E50; GB:M57627; NID:g186270; A;Cross-references: UNIPROT:P22301; UNIPARC:UPI00000034E50; GB:M57627; NID:g186270; A;Cross-references: UNIPROT:P22301; UNIPARC:UPI0000034E50; GB:M57627; NID:g186270; A;Cross-references: UNIPROT:P22301; UNIPARC:UPI0000034E50; GB:M57627; UNIPARC:UPI0000034E50; GB:M57627; UNIPARC:UPI0000034E50; GB:M57627; UNIPARC:UPI0000034E50; GB:M57627; UNIPARC:UPI0000034E50; GB:M57627; UNIPARC:UPI00000034E50; GB:M57627; UNIPARC:UPI00000034E50; GB:M57627; UNIPARC:UPI00000034E50; GB:M57627; UNIPARC:UPI00000034E50; GB:M57627; UNIPARC:UPI000000034E50; GB:M57627; UNIPARC:UPI000000034E50; UNIPARC:UPI000000000000000000000000000000000000</vie></vie></vie>	eb-1992 #text N.; Johnson, , 1991 cytokine synt		123153 128634 AF2545 F81833 E81197 E74651
	NID:g452395; PIDN:CAA51942.1; i	by oligo-cassette PCR walkı	cer. Characterization of the	EMBL:U16720; NID:g1041812; PIDN:AAA80104.1; Straub, H.; Hafner, M.; Tesch, H. 1994	e analysis of human and murintein	GB:M57627; NID:g186270; PIDt rkin, J.; Baldwin, S.; Paliw.	ge Ka	(CGTP) · TI-10	nypothetical prote variant-specific s hypothetical prote conserved hypothet hypothetical prote conserved hypothet hypothetical prote hypothetical prote hypothetical prote

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interleukin 10 - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Species: Sus scrofa domestica (domestic pig)
C;Species: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 09-Jul-2004
C;Accession: I46591
R;Blancho, G.; Gianello, P.; Germana, S.; Baetscher, M.; Sachs, D.H.; LeGuern, Proc. Natl. Acad. Sci. U.S.A. 92, 2800-2804, 1995
A;Title: Molecular identification of porcine interleukin 10: regulation of expl A;Reference number: I46591; MUID:95224028; PMID:7708727
A;Accession: I46591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: DNA sequence and expression of the B95-8 Epstein-Barr virus A;Reference number: A03794; MUID:84270667; PMID:6087149
A;Contents: annotation; protein coding region
C;Superfamily: interlemkin-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-170 <BAN>
A; Residues: 1-170 <BAN>
A; Cross-references: UNIPROT: P03180; UNIPARC: UPI00000CDBB;
A; Cross-references: UNIPROT: P03180; UNIPARC: UPI00000CDBB;
R; Bankter, A.T.; Biggin, M.D.; Deininger, P.L.; F
R; Baer, R.; Bankter, A.T.; Biggin, M.D.; Deininger, P.L.; F
Nature 310, 207-211, 1984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BCRF1 protein - human herpesvirus 4 (strain B95-8) C;Species: human herpesvirus 4, Epstein-Barr virus C;Date: 25-Feb-1985 #sequence_revision 25-Feb-1985 C;Accession: A03741; S32974 R;Bankier, A.T.; Deininger, P.L.; Farrell, P.J.; E
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A;Title: Sequence analysis of the 17,166 bp EcoRI fragment A;Reference number: A93065; MUID:85035713; PMID:6092825
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A;Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STGLKTLNLGSCV----IATNIQEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QTPDHYTLRKISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAV 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LQEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YLGCQALSEMIQFYLEEVMPQAENQDPDIKAHVNSLGENLKTLRLRRCHRFLPC---E
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                                                                                                                                                                                                                                                                                                                                  YKAMSEFDIFINYIE
                                                                                                                                                                                                                                                                                                                                                                              VKALGELDILLOWME 173
                                                                                                                                                                                                                                                                                                                                                                                                                               ENODPEAKDHVNSLGENLKTLRLRLRRCHRFLPC---ENKSKAVEQIKNAFNKLQ-EKGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LRDLRDAFSRVKTFFQTKD-EVD-NLLLKESLLEDFKGYLGCQALSEMIQFYLEEVMPQA
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27.4%;
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28.9%;
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evision 25-Feb-1985
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Pred. No. 8.1e-08;
2; Mismatches 71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 155; DB 1;
Pred. No. 1.1e-07;
9; Mismatches 61
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                                                                           expression
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    RESULT 5
A34853
interleukin-10
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interleukin 10 homolog - equine herpesvirus 2

N;Alternate names: hypothetical protein E7

C;Species: equine herpesvirus 2

C;Date: 17-Feb-1994 #sequence_revision 17-Feb-1994 #text_change 09-Jul-2004

C;Accession: A48558; S55664

R;Rode, H.J; Janssen, W.; Rosen-Wolff, A.; Bugert, J.J.; Thein, P.; Becker, Virus Genes 7, 111-16, 1993

N;Title: The genome of equine herpesvirus type 2 harbors an interleukin 10 (1 A;Reference number: A48558; MUID:93227565; PMID:8385838

A;Accession: A48558
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A; Residues: 1-175 <BLA>
A; Cross-references: UNIPROT: Q29055;
C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q89451; UNIPARC:UPI000012D058; GB:S59624; NID:g299468; A;Note: sequence extracted from NCBI backbone (NCBIN:129334, NCBID:129335)
A;Note: sequence extracted from NCBI backbone (NCBIN:129334, NCBID:129335)
A;Telford, E.A.R.; Watson, M.S.; Aird, H.C.; Perry, J.; Davison, A.J.
J. Mol. Biol. 249, 520-528, 1995
A;Title: The DNA sequence of equine herpesvirus 2.
A;Reference number: S55594; MUID:95302501; PMID:7783207
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C; Superfamily:
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A; Residues: 1-179 < TEL>
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A; Residues: 1-179 < ROD>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   116 FLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWME
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                                                                                      97
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                                                                                                                                                                                                                 39 LQEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNY
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                                         AAVVKALGELDILLQWME
                                                                                      QAENHSTDQEKDKVNSLGEKLKTLRVRLRRCHRFLPC---ENKSKAVEQVKSAFSKLQ-E
                                                                                                                                                                              LHELRAAFSRVKTFFQMKD-QLD-NMLLDGSLLEDFKGYLGCQALSEMIQFYLEEVMP--
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KGVYKAMSEFDIFINYIE 170
                                                                                                                                  QTPDHYTLR---KISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQ 155
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                                                                                                                                                                                                                                                                  24;
                                            173
                                                                                                                                                                                                                                                                  Score 147.5; DB 2
Pred. No. 6.1e-07;
4; Mismatches 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 155;
Pred. No. 1
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precursor -

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R;Feng, L.; Tang, W.W.; Chang, J.C.C.; Wilson, C.B.
Biochem. Biophys. Res. Commun. 192, 452-458, 1992
A;Title: Molecular cloning of rat cytokine synthesis inhibito
A;Reference number: JN0475
A;Accession: JN0475
A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 1-178 cFEN-
A;Cross-references: UNIPROT:P29456; UNIPARC:UPI0000167959
A;Cross-references: UNIPROT:P29456; UNIPARC:UPI0000167959
A;Goodman, R.E.; Oblak, J.; Bell, R.G.
Biochem. Biophys. Res. Commun. 189, 1-7, 1992
A;Title: Synthesis and characterization of rat interleukin-10
A;Reference number: JC1357; MUID:93080542; PMID:1280414
A;Accession: JC1357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N;Alternate names: cytokine synthesis inhibitory factor C;Species: Rattus norvegicus (Norway rat) C;Date: 30-Sep-1993 #sequence revision 30-Sep-1993 #text_change C;Accession: JN0475; JC1357; S36021
                                                                                  A;Molecule type: mRNA
A;Residues: 1,'P',3-64,'L',66-178 <GOO>
A;Cross-references: UNIPARC:UPI000012D477; GB:L02926;
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C;Superfamily: interleukin-10
C;Keywords: cytokine; lymphokine; T-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P18893; UNIPARC:UPI000002149E; R;Kim, J.M.; Brannan, C.I.; Copeland, N.G.; Jenkins, N.A.; J. Immunol. 148, 3618-3623, 1992
A;Title: Structure of the mouse IL-10 gene and chromosomal A;Reference number: I56136; MUID:92268508; PMID:1350294
A;Accession: I56136
                   submitted to the EMBL Data Library, July 1991 A,Reference number: S36021
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Moore, K.W.; Vieira, P.; Florentino, D.F.; Trounstine, M
Science 248, 1230-1234, 1990
A;Title: Homology of cytokine synthesis inhibitory factor
A;Reference number: A34853; MUID:90273182; PMID:2161559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-178 < R
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Best Local S
Matches 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LLELRTAFSQVKTFFQTKD-QLD-NILLTDSLMQDFKGYLGCQALSEMIQFYLVEVMPQA
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31.1%; Pred. No. 1.
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                                                                                       NID:g204903; PIDN:AAA41425.1;
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C; Keywords:
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A;Introns: 1/3; 38/3; 81/1; 143/1; 211/1; 238/2; 278/3; 300/3; 336/2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-567 <SON>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Song, H.J.; Rossi, A.; Ceci, R.; Kim, I.G.; Anzano, M.A.; Jang, S.I.; De Laurenzi, V Biochem. Biophys. Res. Commun. 235, 10-14, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Homo sapiens (man)
C;Date: 02-Sep-1997 #sequence
C;Accession: JC5538
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A; Residues: 1-178 < FE2>
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                                                                                                                                                                                                                                  1 MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQE--IRNGFSD---IRGSVQA
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                             EPQAAVVKAL
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PPALAALRCL
                                                                                                                                 EKSTVLQSELESCKELQELEPENKWCLLTI ILLMRALDPLLYEKETLQYFQTLKAVDPMR
                                                                                                                                                               KDGNIDIRILRRTESLODTKPANRCCLLRHLL-----RLYLDRVFKNYQT-----PDH 103
                                                               ATYLDDLRSKFLLENSVLKMEYAEVRVLHLAHKDLTVLCHLEQLLLVTHLDLSHNRLRTL
                                                                                              YT----LRKISSLANSFLTIK----KDLRLCHAHMTCHCGEEAMKKYSQI-LSH--FEKL 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HLNSLGEKLKTLWIQLRRCHRFLPC---ENKSKAVEQVKNDFNKLQDK-GVYKAMNEFDI
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488
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Pred.
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Pred. No. 8e-06;
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                                                                                                                                                                                                                                                                                   94.5;
No. 0.
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PMID:9196026
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41

RESULT

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A45977
Rab geranylgeranyl transferase component B alpha subunit - rat C:Species: Rattus norvegicus (Norway rat)
C:Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A45977
R;Armstrong, S. A.; Seabra, M.C.; Sudhof, T.C.; Goldstein, J.L.; Brown, M.S.
J. Biol. Chem. 268, 12221-12229, 1993
J. Hiel: cDNA cloning and expression of the alpha and beta subunits of rat Rab. A;Reference number: A45977; MUID:93280201; PMID:8505342
A;Accession: A49977
A;Bolecule type: mRNA; protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probable beta-adaptin YKS5 - yeast (Saccharomyces cerevisiae)

(),Alternate names: protein G9331; protein YGR261c

C;Species: Saccharomyces cerevisiae

C;Date: 17-May-1996 #sequence_revision 17-May-1996 #text_change 09-Jul-2004

C;Accession: S64594; S63450

C;Paccession: S64594; S63450

R;Panzeri, L.; Agostoni Carbone, M.L.; Melchioretto, P.; Plevani, P.; Martegani, submitted to the Protein Sequence Database, May 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT
S64594
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A; Residues: 1-26, 'PLSWVNPP', 36-723, 'T', 725-795, 801-809 .
A; Cross-references: UNIPARC: UPI0000168A41; EMBL: U35411;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Reference number: S63439
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A; Residues: 1-809 < PAN>
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A; Accession: S64594
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: S63450
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                                                                                                                                                                                                                                                                    ;Map
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cross-references: UNIPROT:P46682; UNIPARC:UPI00000530A8; Experimental source: strain S288C; Experimental source: strain S288C; Robinson, L.C.; Engle, H.M.; Pannek, H.R. ubmitted to the EMBL Data Library, September 1995; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 1; Description: Suppressors of loss of yeast casein kinase 2; Description: Suppressors of loss of yeast casein kinase 2; Description: Suppressors of loss of yeast casein kinase 2; Description: Suppressors of loss of yeast casein kinase 2; Description: Suppressors of loss of yeast casein kinase 2; Description: Suppressors of loss of yeast casein kinase 2; Description: Suppressors of loss of yeast casein kinase 3; Description: Suppressors of loss of yeast casein kinase 3; Description: Suppressors of loss of yeast casein kinase 3; Description: Suppressors of loss of yeast casein kinase 3; Description: Suppressors of loss of yeast casein kinase 
                                                                                                                                                                                                                                                                                                                                 Gene: SGD:YKS5
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Best Local (
                                                                                                                                                                Query Match
Best Local Similarity
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EAAAVATSKLGESSYTYY--SQNINPOOLVTLLNSRNSREVRDAMKRIISIMASDDDSID
                                                                 KASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNID
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                                                                                                                                                                Score 93.5; DB Pred. No. 0.66;
                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8O;
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                                                                                                                                  39;
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RESULT 10
F83128
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                                                                                                                                                                                                                                                                                               R;Johnson, D.; Biewaid, i.
submitted to the EMBL Data Library, July 1998
submitted to the EMBL Data Library, July 1998
cosmid Cl6All.
                                                                                                                                                                                                                                                                                                                                                                                         hypothetical protein C16A11.5 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: DNA
A;Residues: 1-140 <STO>
A;Cross-references: UNIPROT:Q9HWP6; UNIPARC:UPI00000C5C2C; GB:AE004830; GB:AE004091; NIC
A;Experimental source: strain PAO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: Complete genome se
A;Reference number: A82950;
A;Accession: F83128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Accession: F83128
R; Stover, C.K.; Pham, X.Q.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probable transcription regulator PA4135 [imported] - Pseudomonas aeruginosa C;Species: Pseudomonas aeruginosa C;Species: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
                                                                             A; Map position: 2
A; Introns: 35/3; 72/1;
                                                                                                                                                                                                                                                                     A; Reference number: A; Accession: T33346
                                                                                                                                                                                                                                                                                          A;Description: The sequence A;Reference number: Z21328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .; Lory, S.; Olson, M.V. Nature 406, 959-964, 200
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                                                                                                                                                                                                                          A; Molecule type: DNA
                                                                                                                                                                                                                                             A;Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local :
                                                                                                                          Gene: CESP:C16A11.5
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Matches
                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 129 DSNSELR-CFA--LSALSDMKMSSLAPIILHTVKKLVTDPSAMVRGEVALAIIKLYRAGK 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        40 IRILRQQGEMESYQLANQACILRPSMTGVLARLERDGIVRRWKAPKDQR-RVYVNLTEKG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62 IRILARTESLODTKPANRCCLLRH-----LLRLYLDRVFKNYQTPDHYTLRKISSLANS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                79 VQLYFADVVKNITTNDTK-----VKRLIHLYLLRFAEN---DPNLTLLSINSLQKSLS 128
l Similarity
36; Conserv
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ilarity 24.5%;
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Conservative
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                                                                               94/3; 129/3; 184/3; 205/1; 279/3; 387/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence of Pseudomonas aeruginosa PAO1, 50; MUID:20437337; PMID:10984043
                   9.0%; Score 81.5;
26.1%; Pred. No. 7.
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22;
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Pred.
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                   5
                                        DB 2;
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49;
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                                        Length 646;
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                                                                                 505/2;
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  31;
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A.; Larbig,
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  Gaps
                                                                                   540/2;
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K.; Li
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protein kinase (
C;Species: Schiz
C;Date: 27-Jan-1
                                 hypothetical protein C44Cl0.4 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change C;Accession: T19925 R;Cottage, A. submitted to the EMBL Data Library, February 1996 A;Reference number: Z19197
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A; Molecule type: DNA
A; Residues: 1-1062 <SAU>
A; Cross references: UNITARC: UPT0000127259; EMBL: AL035537;
A; Cross references: strain 972h-; cosmid c21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P41892; UNIPARC:UPI0000127259; EMBL:X78799; NID:g521098; A;Experimental source: wildtype 972 h minus
A;Note: mRNA sequencing was also done to confirm the intron borders
R;Saunders, D.; Harris, D.; Wood, V.; Rajandream, M.A.; Barrell, B.G.
submitted to the EMBL Data Library, February 1999
A;Reference number: Z21860
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S46367
A;Reference number:
A;Accession: T19925
A;Status: preliminar
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A;Introns: 9/3; 54/3; 64/3; 86/2; 171/1; 325/1; 405/3; 427/1; 544/3; 589/3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Fankhauser, C.; Simanis, V.
EMBO J. 13, 3011-3019, 1994
A;Title: The cdc7 protein kinase is a dosage dependent A;Reference number: S46367; MUID:94313982; PMID:8039497
A;Accession: S46367
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C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change 05-Oct-2004
C;Accession: S46367; T19888
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                                                                                                                                                                                                               RESULT 13
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A; Residues: 1-1062 < FAN>
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Best Local
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preliminary; translated from
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2-isopropylmalate synthase, probable (leuA-1) [imported] - Sulfolobus solfataricus C;Species: Sulfolobus solfataricus C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 09-Jul-2004 C;Accession: B90153
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A;Title: The Genome of the Natural
A;Reference number: AB2577; MUID:21
A;Accession: AD3160
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A;Residues: 1-542 <WIL>
A;Residues: 1-542 <WIL>
A;Cross-references: UNIPROT:Q18616; UNIPARC:UPI000017B837;
A;Experimental source: clone C44Cl0
C;Genetics:
A;Gene: CESP:C44Cl0.4
A;Map position: X
A;Introns: 66/3; 229/3; 309/1; 328/3; 398/3
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A;Experimental source: strain C58 (Dupont)
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A; Residues: 1-502 < KUR>
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294, 2317-2323, 2001
rs: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan,
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R;She, Q.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Chan-Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, F. arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001
A; Jescription: Sulfolobus solfataricus complete genome.
A;Reference number: A99139
A;Acteus; preliminary
A;Acteus; preliminary
A;Rolecule type: DNA
A;Residues: 1-553 <KUR>
A;Cross-references: UNIPROT:Q980Z1; UNIPARC:UPI000006418D; GB:AE006641; NID:g13813259; F. C;Genetics:
A;Gene: leuA-1
C;Superfamily: 2-isopropylmalate synthase leuA
Search completed: December 24, 2005, 10:48:30 Job time : 39 secs
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ata; Euteleostomi; Catarrhini; Hominidae; C.L., Vi Q., Brender T., Haugen H., West J., West J., Chandrasekher Y.A.; Chandrasekher Y.A.; Chandrasekher Y.A.; Chandrasekher Y.A.; West J., Chandrasekher Y.A.; Chandrasekher Y.A.; West J., Chandrasekher Y.A.; West J., Brush J., Wagts A., Xie MH., Yagts A., Xie MH., Goddard A.D., Stinson J., Vagts A., Xie MH., Goddard A.D., Stinson J., Goddard A.D., Stinson J., Goddard A.D., Stinson J., Schuler G.D., Schuler G.D., Schuler G.D., Bhat N.K.,	Q9tsj4 oryctolagus Q9tvd3 oryctolagus P68678 equine herp P68677 equine herp P68677 equine musculu Q6fg99 homo sapien Q923t1 sigmodon hi Q97798 trichosurus Q28474 equus cabal Q6vv71 bubalus bub Q9G311 cercopithec P43480 bos taurus Q8mkg9 saimiri sci P48411 canis famil

WA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

AD Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

AN Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

AN Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

AN Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

AN Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

AN Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

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Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

AN Whiting M., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G.,

AN Holton E., Krimwood J., Schmutz J., Myers R.M.,

AN Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

AN Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

AN Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

AN Blakesley R.W., Touchman J.E., Jones S.J.M., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human

PROTEIN SEQUENCE OF 25-39.
PtbMed=15340161; DOI=10.1110/ps.04682504;
Zwang Z., Henzel W.J.;
"Signal peptide prediction based on anning."

and mouse cDNA sequences.";

Acad.

U.S.A.

99:16899-16903 (2002)

human

verified

peptide prediction d cleavage sites.";

Protein Sci. 13:2819-2824(2004).
-!- FUNCTION: Cytokine that may be psoriasis. Acts through STAT3.
-!- SUBCELLULAR LOCATION: Secreted -!- TISSUE SPECIFICITY: Expressed and other tissues.

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SIMILARITY: Belongs to the IL-10 family.

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Ensembl; ENSG00000162891; Homo sapiens.

R HGNC; HGNC;6002; IL20.

R GO; GO:0005576; C:extracellular region; TAS.

GO; GO:0005577; F:interleukin-20 receptor binding; TAS.

GO; GO:0045617; P:positive regulation of epidermal cel

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GO; GO:0045617; P:positive regulation of tyrosine phos

GO; GO:0045517; P:positive regulation of tyrosine phos

GO; GO:0045517; P:positive regulation of tyrosine phos

GO; GO:0045517; P:positive regulation of tyrosine phos

GO; GO:0050727; P:regulation of inflammatory response;

InterPro; IPR012351; Cytokine 4 hlx.

InterPro; IPR012352; IL-10 add helix.

InterPro; IPR000098; InterTeukin_10.

Pfam; PF00726; IL10; 1.

PFODOm; PD003687; InterLeukin_10.

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tyrosine phosphoryla.
                                                                                                                                                                                                                       epidermal cell
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Best Local S
Matches 175
CONFLICT
CONFLICT
SEQUENCE
                                                 EMBL; A
HSSP; Q
Ensembl
MGI; MG
GO; GO:
                                                                                                                                                                                                                                                                                                  MEDLINE=21097717; PubMed=11163236; DOI=10.1016/S0092-8674(01)00187-8; Blumberg H., Conklin D., Xu W.F., Grossmann A., Brender T., Carollo S., Eagan M., Foster D., Haldeman B.A., Hammond A., Haugen H., Jelinek L., Kelly J.D., Madden K., Maurer M.F., Parrish-Novak J., Prunkard D., Sexson S., Sprecher C., Waggie K., West J., Whitmore T.E., Yao L., Kuechle M.K., Dale B.A., Chandrasekher Y.A.; "Interleukin 20: discovery, receptor identification, and role in
                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Interleukin-20 precursor (IL-20) (Four alpha helix cytokine ZCYTO10).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IL20 MOI
                    InterPro; IPR012351;
InterPro; IPR000098;
Pfam; PF00726; IL10;
                                                                                                                                                                                                This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=Il20; Synonyms=Zcyto10; Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                epidermal function.";
Cell 104:9-19(2001).
                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                              removed
                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Skin
                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                     Muroidea; Muridae;
                                                                                                                                                                                                                                         psoriasis. Acts through STAT3.
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the IL-10
                                                                                                                                                                                        96
                                                                                                                                                                                                                                                                      FUNCTION: Cytokine that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOUSE
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                                                                                                                                                         AF224267;
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                                                                                                                                                                                      long
                                                                                                                                     ; ENSMUSG00000026416; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQMMEETE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DIRITRITESTODIKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANGFLTIK
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126
20072
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Euarchontoglires; Glires;
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STAT3
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Pred. No. 1.8e-80;
1; Mismatches 0
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8385992500B6C447
                                                                                                                                                          mRNA.
                                                                                                                                      musculus.
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lires; Rodentia;
                                                                                                                                                                                        modified
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Best Local S
Matches 131
                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.

MEDLINE=22257669; PubMed=12370360;

Liao Y.-C., Liang W.G., Chen F.W., Hsu

"IL-19 induces production of IL-6 and
apoptosis through TNF-alpha.";
J. Immunol. 169:4288-4297(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q9UHDO; Q96QR4; Q9NUAO;
28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last anoctation update)
11-MAY-2005 (Rel. 47, Last anoctation update)
11-19 precursor (II-19) (Melanoma differentiation protein-like protein) (NG.1).
Name=IL19; Synonyms=ZMDA1;
Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleo Bukaryota; Metazoa; Euarchontoglires; Primates; Catarrhini;
                                                                                                Kotenko S.
Submitted
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CHAIN
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DISULFID
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PROSITE; Cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
MEDILINE=21040165; PubMed=11196675, DOI=10.1038/sj.gene.6363714;
MEDILINE=21040165; PubMed=11196675, DOI=10.1038/sj.gene.6363714;
Milagher G., Dickensheets H., Eskdale J., Izotova L.S.,
Mirochnitchenko O.V., Peat J.D., Vasquez N., Pestka S., Donnell
                  NUCLEOTIDE SEQUENCE [GENOMIC DNA], Rieder M.J., Carrington D.P., Chung Nickerson D.A.;
                                                                                                                                                                                                                    Sexson S.,
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                                                                                                                                       NUCLEOTIDE
                                                                                                                                                                                                                                      Conklin
                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                     "Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                      "Cloning, expression and initial characterization (IL-19), a novel homolog of human interleukin-10 of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
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n S., Smith D., Lok S.,
sapiens homolog of mel
tted (OCT-1999) to the
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                                                                                                .V., Pestka
(JUN-2001)
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                                                                                                                                   SEQUENCE.
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176
126
132
  HL66682
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                                                                                              to ;
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75.3%;
                                                                                              the
                                                                                                                                                                          Loften-Day C., Whitmore T
S., Pownder T., O'Hara P.
melanoma differentiation
the EMBL/GenBank/DDBJ data
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program
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Pred. No. 1
                                     Chung
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                                   AND VARIANT
g M.-W., Lee
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  for
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TNF-alpha and
  genomic
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                                   PHE-175.
                                                                                                databases
applications,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IL-10).";
                                                                                                                                                                                                 associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of interleukin-19
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                                     C.L.,
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유성

MKLQCVSLWLLGTILILCSVDNHGLR---

-RCLISTDMHHIEESFQEIKRAIQAKDTFP

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MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI

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Query Match
Best Local S
Matches 69
                                                                                                                                                        GO; GO:0005576; C:extracellular region; N
GO; GO:0005125; F:cytokine activity; TAS.
GO; GO:0006955; P:immune response; NAS.
GO; GO:0007165; P:signal transduction; NA
InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR01098; Interleukin_10.
Pfam; PF00726; ILIO; 1.
SMART; SM00188; ILIO; 1.
SMART; SM00188; ILIO; 1.
PROSITE; PS00520; INTERLEUKIN_10; 1.
                                                                                                                            Glycoprotein;
SIGNAL
CHAIN 2
                                                            CARBOHYD
CARBOHYD
DISULFID
DISULFID
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VARIANT
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GO; GO:000
GO; GO:000
GO; GO:000
GO; GO:000
InterPro;
InterPro;
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                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                EMBL;
                                                                                                                                                                                                                                                                                                                                                     EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     helical cytokines.";
J. Biol. Chem. 278:3308-3313(2003).
-!- FUNCTION: May play some important roles
Up-regulates IL-6 and TWF-alpha and indu
similarity).
                                                                                                                                                                                                                                                                                                                                                                                                       This Swiss-Prot entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FHCRC, Seattle, WA (URL: http://pga.gs.washington.edu)."; Submitted (JUN-2001) to the EMBL/GenBank/DDBJ databases. [6]
                                                                                                                                                                                                                                                                                                                                                                                               the European
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PubMed=12403790; DOI=10.1074/jbc.M208602200;
Chang C., Magracheva E., Kozlov S., Fong S.,
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Human chromosome 1 i
Submitted (FEB-2000)
                                                                                                                                                                                                                                                                             HGNC; HGNC:5990;
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Ensembl; ENSG00000142224; H
                                                                                                                                                                                                                                                                                                                                                                                     removed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBCELLULAR SIMILARITY:
                                                                                                                                                                                                                                                                                                        AF276915; AAG16755.1; -; Genomic_DNA.
AF433946; AAN40906.1; -; mRNA.
AF192498; AAF06663.1; -; mRNA.
AY040367; AAK91776.1; ALT_INIT; mRNA.
AF390905; AAK64498.1; -; Genomic_DNA.
AL049615; CAB72342.1; -; Genomic_DNA.
 69;
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           Similarity
                                         177
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                                                                                                                                               Apoptosis; Cytokine;
Polymorphism; Signal
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/FTId=VAR_013077.
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Pred. No. 9.4
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N-linked (GlcNAc.
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QSVUT3 HUMAN
QSVUT3 HUMAN PRELIMINARY;
AC QSVUT3;
AC QSVUT4.29,
AC QSVUT4.29,
AC QSVUT5.29,
AC Interleukin 19.
AC ENKARYOTA; MATAZOA; Chordat.
AC Mammalia; Eutheria; Euarcho.
AC NCBI_TAXID=9606;
AN (1)
AC SUBCELLULAR LOCATION: S.
AC SUBCELLULAR LOCATI
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Best Local S
Matches 69
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13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
Interleukin 10 family protein.
Brachydanio rerio (Zebrafish) (Danio rerio)
                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Bu
Actinopterygii; Neopterygii; Teleostei; Ostariophysi;
Cyprinidae; Danio.
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Pfam; PF00726; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; UNKNOWN_1.
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EMBL; AL513315; CAH71814.1; -; Genomic_DNA.
SMR; QSVUT3; 60-215.
                                                                          NCBI_TaxID=7955
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         NUCLEOTIDE SEQUENCE.
Ram S., Daisuke I.,
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BRARE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          121
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            Sakai M.;
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k; Pred. No. 1.2e
36; Mismatches
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Last sequence update)
Last annotation updat
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SIGNAL
                                                                                                                                                                                                                                                  Ensembl; ENSMUSG00000016524; Mus MGI; MGI:1890472; Il19.
GO; GO:0006917; P:induction of ap GO; GO:0042226; P:interleukin-6 b GO; GO:0006800; P:oxygen and reac GO; GO:0006800; P:oxygen and reac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=Swiss Webster / NIH;
MEDLINE=22257669; PubMed=12370360;
Liao Y.-C., Liang W.G., Chen F.W., Hsu
"IL-19 induces production of IL-6 and
apoptosis through TNF-alpha.";
J. Immunol. 169:4288-4297(2002).
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10-MAY-2005 (Rel. 47, Created)
10-MAY-2005 (Rel. 47, Last sequence upon
10-MAY-2005 (Rel. 47, Last annotation)
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Q8CJ70;
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SEQUENCE 168 AA; 19110 MW; 807D18AD736C709B CRC64;
                                                                                                                                                                     GO; GO:0006917; P:induction of apoptosis; IDA.
GO; GO:0042226; P:interleukin-6 biosynthesis;
GO; GO:0006800; P:oxygen and reactive oxygen s
InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR000098; Interleukin_10.
ProDom; PD003687; Interleukin_10; 1.
                                                                                                                                                                                                                                                                                                                                                                      EMBL; AF453945; AAN40905.1; HSSP; Q9UHD0; 1N1F. SMR; Q8CJ70; 23-170.
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Mammalia; Eutheria;
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-i- SIMILARITY: Belongs to the IL-10 family.
                                                                                                                                     PROSITE; PS00520;
                                                                                                                                                              SMART;
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                                                                                                       INTERLEUKIN 10; FALSE NEG. ne; Glycoprotein; Signal.
                 24
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                                                                                    Glycoprotein; Sig
24 Potential
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No. 1.1e-21;
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TNF-alpha and
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Matches 57
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L BMC Genomics 4:29-29(2003).

EMBL; AY294557; AAP57416.1; -; Genomic_DNA.

R EMBL; AY294556; Caxtracellular space; IEA.

GO; GO:0005615; C:extracellular space; IEA.

GO; GO:0005125; F:cytokine activity; IEA.

GO; GO:0005125; F:cytokine activity: IEA.

R GO; GO:0005125; P:immune response; IEA.

R GO; GO:0005125; F:cytokine 4hlx.

InterPro; IPR000098; Interleukin_10.

P Deam; PF00726; ILI0; 1.
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01-OCT-2003
01-FEB-2005
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Bukaryota; Metazoa; Chordata; Craniata; Verte
Actinopterygii; Neopterygii; Teleostei; Eutel
Acanthomorpha; Acanthopterygii; Percomorpha;
Tetraodontoidea; Tetraodontidae; Tetraodon.
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Q7SX60;
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Mogensen K., Monneron D.;
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Pred. No. 8.4e-18;
3; Mismatches 53;
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Euteleostei; Neoteleostei;
rpha; Tetraodontiformes;
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RESULT 9

Q92584 MUSE

ID Q92584 M

AC Q92584 F

DT 01-DEC-2

DT 01-DEC-2

DT 01-DEC-2

DT 01-MAR-2

DE Melanoma

GN Name=112

OS Nume=12

OC Eukaryot

OC Mammalia

OC Muridae;

OX NCBI Taa

RN 11

RP NUCLEOTI

RA Madiredc

RL Submitte
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Best Local S
Matches 56
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GO; GO:0005615; C:extracellular space; TAS. InterPero; IPR000098; Interleukin 10. ProDom; DD003687; Interleukin 10; 1. SMART; SM00188; IL10; 1.

PROSITE; PS00520; INTERLEUKIN 10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-DEC-2001 (TrEMBLrel. 19, (
01-MAR-2004 (TrEMBLrel. 26, L
Th2-specific cytokine FISP.
Name=1124;
  NUCLEOTIDE SEQUENCE.
STRAIN=129/SvJ;
Madireddi M.T., Lin J
Submitted (FEB-2000)
                                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytokine.
SEQUENCE
                                                                                                                                                                                                                                                             01-DEC-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
01-MAR-2004 (TrEMBLrel. 2
Melanoma differentiation
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Mammalia; Eutheria; Eu
Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                    Q925S4_MOUSE PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     secreted by Th2 cells.";
J. Immunol. 166:5859-5863(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schaefer G., Venkataraman C., Schindle "FISP (IL-4-induced secreted protein),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (1)
                                                                                                                   NCBI_TaxID=10090;
                                                                                                                                                                                                                                               Name=1124;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=21240641; PubMed=11342597;
Schaefer G., Venkataraman C., Schi
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    J.,
                                                                                                                                                                     Chordata; Craniata; Vertebrata; Eute
Euarchontoglires; Glires; Rodentia;
                                                                                                                                                                                             Chordata;
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Euarchontoglires;
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26,
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    Z., Shay J.W., Huberman E.,
EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                    Last sequence update)
Last annotation update)
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Last
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Pred. No. 2.6e
30; Mismatches
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                                                                                                                                                                                             Euteleostomi;
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                             Fisher
                                                                                                                                                                     Sciurognathi;
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Matches
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Q13007; Q950B0; Q95KG4;
Q13007; Q950B0; Q95KG4;
Q1-NOV-1997 (Rel. 35, Created)
Q1-NOV-1997 (Rel. 35, Last sequence update)
Q1-NOV-1997 (Rel. 47, Last annotation update)
Interleukin-24 precursor (Suppression of tumorigenicity (Melanoma differentiation-associated protein 7) (MDA-7).
(Melanoma differentiation-associated protein 7) (MDA-7).
                                                                                    "Genomic structure, chromosomal localization and expression o melanoma differentiation associated gene-7 (mda-7): potential relationship with cellular senescence."; Submitted (FEB-2000) to the EMBL/Garn--7,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HUMAN
                                                                                                                                                                                                                                                                                          MEDLINE=96132699; PubMed=8545104;
Jiang H., Lin J.J., Su Z.-Z., Goldstein N.I., Fisher P.B.;
"Subtraction hybridization identifies a novel melanoma differentiation
associated gene, mda-7, modulated during human melanoma
differentiation, growth and progression.";
Oncogene 11:2477-2486(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ensembl; ENSMUSG00000026420; Mus musculus. MGI; MGI:2135548; Il24. GO; GO:0005615; C:extracellular space; TAS
                                                                                                                                                                                                                    Peat J., Kube D., Eskdale J., Jueliger S., "The human MDA-7 gene.";
Submitted (JUN-2000) to the EMBL/GenBank/DI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR000098; Interleukin_
ProDom; PD003687; Interleukin_10;
SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -!- SUBCELLULAR LOCATION: Secreted (By similarity).
EMBL; AF235006; AAKS2590.1; -; mRNA.
 Rieder M.J., Armel T.Z., Carrington D.P., Chung M.-W., Lee K.I
Poel C.L., Toth E.J., Yi Q., Nickerson D.A.;
"SeattleSNPs. NHLBI HL66682 program for genomic applications,
FHCRC, Seattle, WA (URL: http://pga.gs.washington.edu).";
                                                                         NUCLEOTIDE
                                                                                                                                                                                         NUCLEOTIDE SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Melanoma;
                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEE 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VSGAESCYLAHSLLKFYLNTVFKNYHSKIAKFKVLRSFSTLANNFIVIMSQLQPSKDNSM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LLWNQVPGLEGQEFRSGSCQVTGVVLPELWEAFWTVKNTVQTQDDITSIRLL-KPQVLRN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Metazoa;
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                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             181 AA;
                            Armel T.Z., (
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                                                                         [GENOMIC
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                                                                                                                                                                                          AND VARIANT HIS-124.
                                                                                                                                                                                                                       to the EMBL/GenBank/DDBJ
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Pred. No. 3
                                                                           AND
                                           D.P.,
                                                                           VARIANTS
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les 71;
                                          Chung
                                                                                                                                                                                                                                                  Gallagher
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                                           M.-W.,
                                                                         HIS-124; ARG-125
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RX MEDLINE-99045696; PubMed-9826712; DOI=10.1073/pnas.95.24.14400;
RA SU Z.-Z., Madireddi M.T., Lin J.J., Young C.S.H., Kitada S.,
Reed J.C., Goldstein N.I., Fisher P.B.;
RA Reed J.C., Soldstein N.I., Fisher P.B.;
RA Reed J.C., Matler Reed J.C., Matler Reed J.C., Ree
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Long L., Wallan K., Farmer A.A., Rubin G.M., Hong L.,
A Hopkins S., Loquellano N.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
A Hillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human and many seconds of the second se
                                                                                               GO; GO:0006915; P:apoptosis; TAS.
InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR010098; Interleukin 10.
ProDom; PD003687; Interleukin 10; 1.
SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; U16261; AAA91780.1; -; mRNA.

EMBL; AF276916; AAG41401.1; -; Genomic_DNA.

EMBL; AF235005; AAK52589.1; -; Genomic_DNA.

EMBL; AY062931; AAL34146.1; -; Genomic_DNA.

EMBL; BC009681; AAH09681.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROTEIN SEQUENCE OF 52-66.
PubMed=15340161; DOI=10.1110/ps.04682504;
Zhang Z., Henzel W.J.;
"Signal peptide prediction based on analy
verified cleavage sites.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Proc.
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                                 PROSITE; PS00520; INTERLEUKIN_10; Apoptosis; Cytokine; Direct prote:
                                                                                                                                                                                                                                                                                 MIM;
                                                                                                                                                                                                                                                                                                                                                   Ensembl; ENSG00000162892; HGNC; HGNC:11346; IL24.
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                                 protein
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Grouse L.H., Derge J.G.,
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SEQUENCE
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CHAIN
CARBOHYD
CARBOHYD
CARBOHYD
                                                                                                                                                                                                                  EMBL;
                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
Kalnine N., Chen X., Rolfe
Koundinya M., Raphael J.,
Phelan M., Farmer A.;
                                                                                                                                                                                                                                       Submitted (MAY-2003) to the
                                                                                                                                                                                                                                                                                                                                  NCBI
                                                                                                                                                                                                                                                                                                                                                      Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Q53XZ7_HUMAN PRELIMINARY;
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                                                                                                                                                                               GO; GO:0006955;
                                                                                                                                                                                                                                                           Phelan M., Farmer A.;
"Cloning of human full-length CDSs in BD
                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                        Interleukin 24.
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                                                                                                                                                                                                                                                                                                                                 TaxID=9606;
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                                                                                                                         Similarity
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                                                                                      LAFSLLSAAFYLLWTPSTGL--KTLNLGSCVIATNL-QEIRNGFSDIRGSVQAKDGNIDI
 KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEE 174
                     RLLQQ-EVLQNVSDAESCYLVHTLLEFYLKTVFKNYHNRTVEVRTLKSFSTLANNFVLIV
                                          RILRRTESLODTKPANRCCLLRHLLRLYLDRVFKNY--QTPDHYTLRKISSLANSFLTIK
                                                                  LGFTL-----LLWSQVSGAQGQEFHFGPCQVKGVVPQKLWEAFWAVKDTMQAQDNITSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMBE 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNY--QTPDHYTLRKISSLANSFLTIK 120
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                                                                                                                                                         AA;
                                                                                                                                                                               P:immune response; IEA.
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126
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                                                                                                                                                                                                                                                                                                                                                     Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                       23.8%;
                                                                                                                                                         23912 MW;
                                                                                                                                                                                                                                                                              Rolfs A., Halleck A., Hines L., Eisenstein
1 J., Moreira D., Kelley T., LaBaer J., Lin
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N-linked (GlcNAG...) (Pote N-linked (GlcNAG...) (Pote N-linked (GlcNAG...) (Pote N-linked (GlcNAG...) (Pote Y-> H (in dbsNP:1150258).

/FTId=VAR 011974.

H -> R (in dbsNP:3093431).
/FTId=VAR 013097.
V-> L (in dbsNP:3093446).
/FTId=VAR 03098.
/FTId=VAR 03098.
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Last annotation update
                                                                                                                                                                                                                                                                                                                                                                                                                        Created)
                                                                                                                        Score 217;
Pred. No. 7
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Pred. No. 7.3e-13
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Q9WVP8 RAT 1
Q9WVP8 RAT 1
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DT 01-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                            CARBOHYD
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                   Cytokine;
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zhang R., Tan Z., Liang P.;
"Identification of a novel ligand-receptor pair constitutively
activated by ras oncogenes.";
J. Biol. Chem. 275:24436-24443(2000).
-!- SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ensembl; ENSRNOGO0000004470; Rattus norvegicus. GO; GO:0048144; P:fibroblast proliferation; TAS. GO; GO:0042060; P:wound healing; TAS. InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR000098; Interleukin 10.
ProDOm; PD0003687; Interleukin 10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Marinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Cytokine-like protein Mob-5 precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOB5
                                                                                                                                                                                                                                                                                                                                                                                         CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
MEDLINE=20387329; PubMed=10825166; DOI=10.1074/jbc.M001958200;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- SIMILARITY: Belongs to the IL-10 family.
                                                                                                                                                         148
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                                                                                                                                                                                     93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AF269251; AAF75553.1; -; mRNA.
                                                                                                                                                                                                                                                                                                       49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              long as its content is in no way modified and this statement
                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                           GPCQVTGVVLPELWEAFWTVKNTVKTQDELTSVRLL-KPQVLQNVSDAESCYLAHSLLKF
                                                                                                                              HRTFKQLDIEVALAKAFGEVDILLAWMQ 178
                                                                                                                                               H--FEKLEPQAAVVKALGELDILLQWME 173
                                                                                                                                                                                                          YLDRVFKNYQTP--DHYTLRKISSLANSFLTIKKDLRLCHAHMTCHCGEBAMKKYSQILS
                                                                                                                                                                                       YLNTVFKNYHSKIVKFKVLKSFSTLANNFLVIMSKLQPSKDNAMLPISDSARRRF--LLY
                                                                                                                                                                                                                                                                       GSC-VIATNIQEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRL
                                                                                                                                                                                                                                                                                                                                                                                       Glycoprotein; Signal.
1 28 Po
29 183 Cy
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                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                           76 N
21096 MW;
                                                                                                                                                                                                                                                                                                                   21.0%;
                                                                                                                                                                                                                                                                                                                   Score 191; DB 1;
Pred. No. 2.2e-10;
                                                                                                                                                                                                                                                                                                                                                           Cytokine-like protein Mob-5.
N-linked (GlcNAc. . .) (Pote;
AF7A11466C491AC4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                      Potential
                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                Length 183;
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                                                                                                                                                                                                                                                                                                                                                                           (Potential).
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(TrEMBLrel. (TrEMBLrel. (TrEMBLrel.

12, 12, 26,

Created)
Last sequence up
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update)

PRELIMINARY;

183

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Q6A2H5
ID Q6A2H5
ID Q6A2H5
DT 255
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Best Local S
Matches 47
                                   "Cloning and Characterization of Chicken II
Immune Response to Eimeria maxima.";
J. Immunol. 173:2675-2682(2004).
EMBL; AJ621254; CAF18432.1; -; mRNA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0005125; F:cytokine activity; IEA.
GO; GO:0006955; P:immune response; IEA.
InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR012351; IL-10 add helix.
InterPro; IPR012352; IL-10 add helix.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C49a.

C49a.

Rattus norvegicus (Rat).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Eukaryota; Butheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;

Mammalia; Eutheria; Bartus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QGA2H5_CHICK PRELIMINARY;
Q6A2H5;
Q6CT-2004 (TremBLrel. 28,
25-QCT-2004 (TremBLrel. 28,
25-QCT-2004 (TremBLrel. 28,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000098; Interleukin 10. ProDom; PD003687; Interleukin 10; 1. PROSITE; PS00520; INTERLEUKIN 10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Archosauria; Aves;
                                                                                                                                                                                                                                                                                                                                                                                 Rothwell L., Young J.,
Archer A., Smith A.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gallus gallus (Chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interleukin-10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 47; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TFKQLDIEVALAKAFGEVDILLAWMQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HFEKLEPQAAVVKALGELDILLOWME 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GPCQVTGVVLPELWEAFWTVKNTVKTQDELTSARLL-KPQVLQNVSDAESCYLAHSLLKF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GSC-VIATNIQEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Chordata; Craniata; Vertebrata; Euteleostomi; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21052 MW; FF69A96AFC473E4F CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                         Kaiser P.;
                                                                                                                                                                                                                                                                                                                                                                                                                     Zoorob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29;
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Last
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Pred. No. 5
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5.5e-10;
66;
                                                                                                                                                                                                                     IEA.
                                                                                                                                                                                                                                                                                                                                                         IL-10
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                                                                                                                                                                                                                                                                                                                                                                                                                         Hesketh P.,
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RESULT 15
Q6A2H4
Q6A2H
Q6A2H
AC Q6A2H
AC Q6A2H
DT 25-00
DT 25-00
DE Inte
GN Name
OS Gall
OC Buke
OC Arch
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OC Arch
CRA NCB
RN NCB
RN NCB
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Best Local S
Matches 53
                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immune Response to Elneria maxima.";

J. Immunol. 173:2675-2682(2004).

EMBL; AJ621614; CAF21727.1; -; Genomic DNA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0005525; F:cytckine activity; IEA.
GO; GO:0006955; P:immune response; IEA.
InterPro; IPR012351; Cytckine 4 hlx.
InterPro; IPR012352; IL-10 add helix.
InterPro; IPR00098; InterTeukIn_10.
Pfam; PF00726; IL10; 1.
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25-OCT-2004 (TrEMBLrel.
25-OCT-2004 (TrEMBLrel.
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Q6A2H4;
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ProDom; PD003687; Interleukin
SMART; SM00188; IL10; 1
                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rothwell L., Young J., Archer A., Smith A.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Archosauria; Aves; Neognathae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Interleukin-10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Cloning and Characterization of Chicken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHICK
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117
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                                                                                                                                                                                                                                                                                      53;
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                                                                                                                                                                                                                                                                                                            Similarity
  KATIRRCHRFFTC--
                                                                                                                                          IDIRILARTESLODTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTI
                                                                                                                                                                                                                                     MKASSLAFSLLSAAFYLLWTPSTGLKTLNLG-SCVIATNLQEIRNGFSDIRGSVQAKDGN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KATMRRCHREFTC---EKRSKAIKQIKETFEKMD-ENGIYKAMGEFDIFINYIEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEE
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                                            KKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEE
                                                                                           T'NI OLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                 AA;
                                                                                           -SSELLDEFKGTFGCQSVSEMLRFYTDEVLPRAMQTSTSHQQSMGDLGNMLLGL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Interleukin_10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20514 MW;
                                                                                                                                                                                                                                                                                                          19.3%;
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Pred. No. 7e-09;
2; Mismatches
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; Galliformes; Phasianidae; Phasianinae;
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Search completed: December 24, 2005, 10:46:54 Job time : 230 secs

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Command line parameters:

-MODEL-frame+ p2n.model -DEV=xlp
-MODEL-frame+ p2n.model -DEV=xlp
-Q-/cgn2 1/USPTO_spool p/US10789129/runat 23122005_152720_20263/app query.fasta_1.327
-Q-/cgn2 1/USPTO_spool p/US10789129/runat_23122005_152720_20263/app query.fasta_1.327
-Q-/cgn2 1/USPTO_spool p/US10789129/cgn2 -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-DB=GENEMB1 -QFMT=fastap -SUPFIX=p2n.rge -MINMATCH=0.1 -LOOPCL=0 -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXIEN=2000000000
-USER-US10789129 @CGN 1 17415 @runat 23122005 152720 20263 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-PGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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069364 Homo sap	BC06936		ø	4	
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Conklin, D.C., Haldeman, B.A. and Grossmann, A.

Mammalian cytokine-like polypeptide-10

Patent: JP 2001524313-A 1 04-DEC-2001;

ZYMOGENETICS INC

OS Homo sapiene (human)

PN JP 2001524313-A/1

PD 04-DEC-2001

PD 04-A61P17/00, PC A61P35/00,A61P37/02,A61P43/00,C07K14/52,C07K16/24,C07K16/42. PC C12N15/00, Homo sapiens (human) BD087297.1 GI:22632907 JP 2001524313-A/1. Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Mammalian cytokine-like polypeptide-10. Hominidae; Homo. 926 bp DNA linear PAT 27-AUG-2002 ဂ္ဂ

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                                        Conklin,D.C., Haldeman,B.A. and Grossmann,A. Mammalian cytokine-like polypeptide-10 Patent: EP 1424393-A 1 02-UUN-2004;
                                Zymogenetics Inc
                                                                                                                               Homo sapiens (human)
                                                                                     Hominidae; Homo.
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Moore,M.D.
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Xu, W., Kindsvogel, W.R., Chen, Z., Hughes, S.D., Chandrasekher, Y.A.,
Dillon, S.R., Lehner, J.M., Siadak, A.W., Sivakumar, P.V. and
                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.
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Mammalian cytokine-like polypeptide-10
Patent: US 6576743-A 1 10-JUN-2003,
ZymoGenetics, Inc.; Seattle, WA
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Blumberg,H., Conklin,D., Xu,W.F., Grossmann,A., Brender,T., Carollo,S., Eagan,M., Foster,D., Haldeman,B.A., Hammond,A., Haugen,H., Jelinek,L., Kelly,J.D., Madden,K., Maurer,M.F., Parrish-Novak,J., Prunkard,D., Sexson,S., Sprecher,C., Waggie,K.,
                                         Hominidae; Homo.

1 (bases 1 to 926)

Blumberg, H., Conklin, D.,
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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GluProGlnAlaAlaValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet
                                          AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                                                                                                                                                                                                                              TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGACATACGGGGCAGT
                                                                                                   GCCAATTCCTTTCTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCCACATGACATGC
                                                                                                                           AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
                                                                                                                                                                          ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
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West,J., Whitmore,T.E., Yao,L.,
Chandrasekher,Y.A.
Interleukin 20: discovery, recep
epidermal function
Cell 104 (1), 9-19 (2001)
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Direct Submission
Submitted (12-JAN-2000) Biomolecular Informatics,
Inc., 1201 Eastlake Ave. East, Seattle, Wa. 98102.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conklin,D., Grossmann,A., Haldeman,B. and O'Hara,P. Homo sapiens four alpha helix cytokine allele 1 Unpublished
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Methods
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Sequence 18 from Patent
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                                      GluGluThrGlu 144
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                                                                                                                      CACTGTGGTGAGGAAGCAATGAAAAAAATACAGCCAGATTCTGAGCCACTTCGAAAAAACTG
                                                                                                                                                                                         AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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Strauberg, R.D., Collins, F.S., Wagner, L., Schamen, C.M., Schuler, G.D.,

Klausner, R.D., Collins, F.S., Wagner, L., Schaefer, C.F., Bhat, N.K.,

Altschul, S.F., Zeeberg, B.F., Buetow, K.H., Schaefer, C.F., Bhat, N.K.,

Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsich, F.,

Diatchenko, L., Warusina, K., Farmer, A.A., Rubin, G.M., Hong, L.,

Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S.,

Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J.,

Abramson, R.D., Mullahy, S.J., Bosak, S.A., McEwan, P.J.,

McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S.M.,

Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hulyk, S.W.,

Villalon, D.K., Muzny, D.M., Sodergren, B.J., Lu, X., Gibbs, R.A.,

Villalon, D.K., Muzny, D.M., Sodergren, B.J., Lu, X., Gibbs, R.A.,

Fahey, J., Helton, B., Ketteman, M., Madan, A., Rodrigues, S.,

Sanchez, A., Whiting, M., Madan, A., Young, A.C., Shevchenko, Y.,

Bouffard, G.G., Blakesley, R.W., Touchman, J.W., Green, B.D.,

Dickson, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M.,

Schnerch, A., Schein, J.E., Jones, S.J., and Marra, M.A.

Mammalian Gene Collection program Team

Generation and initial analysis of more than 15,000 full-length

human and mouse cDNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tissue Procurement: Baylor Human Genome Sequencing Center cDNA Library Preparation: Baylor Human Genome Sequencing CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL) DNA Sequencing by: Baylor College of Medicine Human Genom Sequencing Center College of Medicine Human Genom Sequencing Center
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Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged, H.,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (29-APR-2004) National Institutes of Health, Mammalian Gene Collection (MGC), Bethesda, MD 20892-2590, USA
NIH-MGC Project URL: http://mgc.nci.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                           DEFINITION
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Hominidae; Homo.

1 (bases 1 to 531)

Strausberg,R.L., Feingold,E.A., Grouse,L.H., Derge,J.G.,
Klausner,R.D., Collins,F.S., Wagner,L., Shenmen,C.M., Schuler,G.D.,
                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                         Homo
                                                                                                                                  BC069364.1 GI:47479475
                                                                                                                                                           IMAGE:7262128), complete
                                                                                                                                                                         Homo sapiens interleukin
                                                                                                                                                                                                                                                                                                                   GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
                                                                                                                                                                                                                                                                                                                                                                          HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                 AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GluGluThrGlu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
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                                                                                                                                                                                                                                             GAGGAGACAGAA
                                                                                                                                                                                                                                                                                                    GAACCTCAGGCAGCAG:
                                                                                                                                                                                                                                                                                                                                                          CATTGTGGGGAAGCAATGAAGAAATACAGCCAGATTCTGAAGTCACTTTGAAAAAGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                 GCCAATTCCTTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGGGTATTTAAAAACTACCAGACCCCTGACCATTATACTCTCCGGAAGATCAGCAGCCTC
                                                                                        sapiens
                                                                                                    sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /product="interleukin 20, product="interleukin 20, protein id="AAH69311.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /translation="mkasslafsllsaafyllwtpstglktlnlgscvlatnlqeirn
Gfseirgsvqakdgnidirilrrteslqdtkpanrccllrhllrlyldrvfknyqtpd
Hytlrkisslansfltikkdlrlchahmtchcgeeamkkysqilshfeklepqaavvk
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/codon_start=1
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one MGC:96919
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PUBMED
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AUTHORS
TITLE
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COMMENT
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Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K., Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F., Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L., Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L., Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S., Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J., Abramson, R.D., Mullahy, S.J., Bosak, S.A., McEwan, P.J., McKernan, K.J., Mullahy, S.J., Bosak, S.A., McEwan, P.J., McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S., Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hulyk, S.W., Villalon, D.K., Muzny, D.M., Sodergren, E.J., Lu, X., Glbbs, R.A., Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S., Sanchez, A., Whiting, M., Madan, A., Young, A.C., Shevchenko, Y., Bouffard, G.G., Blakesley, R.W., Touchman, J.W., Green, B.D., Dickson, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M., Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E., Schnerch, A., Schein, J.E., Jones, S.J. and Marra, M.A. Generation and initial analysis of more than 15,000 full-length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyl
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzi
A.N., Gibbs, R.A.
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Tissue Procurement: Baylor Human Genome Sequencing Center
CDNA Library Preparation: Baylor Human Genome Sequencing Center
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Baylor College of Medicine Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clone distribution: MGC clone distribution information can be for through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRBR Plate: 1 Row: d Column: 2.
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Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
Submitted (29-APR-2004) National Institutes of Health, Mammalian Gene Collection (RGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Center code: BCM-HGSC
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DB:
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RS Strausberg, L., Feingold, E.A., Grouse, L.H., Derge, J.G.,
RS Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D.,
Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K.,
Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F.,
Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L.,
Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L.,
Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S.,
Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J.,
Abramson, R.D., Mullahy, S.J., Mouellano, N.A., Peters, G.J.,
McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S.,
Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hilyk, S.W.,
Villalon, D.K., Muzny, D.M., Sodergren, E.J., Lu, X., Gibbs, R.A.,
Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S.,
Sanchez, A., Whiting, M., Madan, A., Young, A.C., Shevchenko, Y.,
Bouffard, G.G., Blakesley, R.W., Touchman, J.W., Green, E.D.,
Dickson, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M.,
Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E.,
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC
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중 음 중

ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40

CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer 20

97

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JOURNAL
PUBMED
REFERENCE
AUTHORS
CONSRIM
TITLE
JOURNAL
                                                          Percent Similarity:
Best Local Similarity:
Query Match:
US-10-789-129-26 (1-144) x BC069398
                                                                                                                              Score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRBR Plate: 1 Row: e Column: 2.
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged, H.,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
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Mammalian Gene Collection Program Team
Generation and initial analysis of more than 15,000 full-length
human and mouse cDNA sequences
Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2002)
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Hytlrkisslansfltikkdlrlchahmtchcgeeamkkysqilshfeklepqaavvk
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                                          4.73e-72
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Baylor Human Genome Sequencing Center
cDNA Library Preparation: Baylor Human Genome Sequencing Center
                                                                                  Direct Submission
Submitted (29-APR-2004) National Institutes of Health, Mammalian Submitted (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                          NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens interleukin IMAGE:7262152), complete
                                                                                                                                          2 (bases 1 to 531)
Director MGC Project.
                                                                                                                                                                                    human and mouse cDNA sequences Proc. Natl. Acad. Sci. U.S.A.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny,
A.N., Gibbs, R.A.
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DNA Sequencing by: Baylor College of Medicine Human Genome
Sequencing Center
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HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
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                                                                                        AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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Hytlrkisslansfltikkdlrlchahmtchcgeeamkkysqilshfeklepoaavvk
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/db xref="GeneID:50604"
/db xref="MIM:605619"
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Mammalian Gene Collection Program Team
Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences

Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Director MGC Project.

Direct Submission

Submitted (29-APR-2004) National Institutes of Health, Mammalian Submitted (29-APR-2004) National Institutes Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                               Clone distribution: MGC clone distribution through the I.M.A.G.E. Consortium/LLNL at: Series: IRBR Plate: 1 Row: g Column: 2.
                                                                                                                                                          Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, (
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny,
A.N., Gibbs, R.A.
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                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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Mammalia; Eutheria;
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IMAGE:7262164), complete
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/organism="Homo sapiens"
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Homo sapiens interleukin IMAGE:7262176), complete BC069487
BC069487.1 GI:47480770
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HYTLRKISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVK
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/protein_id="A4H6949.1"
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/tissue_type="PCR resqued clones"
/clone_lib="NIH_MGC_244"
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Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Collins FS, Wagner L, Shemmen CM, Schuler GD, Altschul SF, Zeeberg B, Buetow KH, Schaefer CF, Bhat NK, Hopkins RP, Jordan H, Moore T, Max SI, Wang J, Hsieh F, Diatchenko L, Marusina K, Farmer AA, Rubin GN, Hong L, Stapleton M, Soares MB, Bonaldo MF, Casavant TL, Scheetz TE, Brownstein MJ, Usdin TB, Toshiyuki S, Carninci P, Prange C, Raha SS, Loquellano NA, Peters GJ, Abramson RD, Mullahy SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SS, MocEwan PJ, McKernan KJ, Gibbs RA, Fahey J, Helton E, Ketteman M, Madan A, Rodrigues S, Sanchez A, Whiting M, Wadan A, Young AC, Shevchenko Y, Bouffard GG, Blakesley RW, Touchman JW, Green ED, Dickson MC, Rodriguez AC, Grimwood J, Schmutz J, Myers RM, Butterfield YS, Krzywinski MI, Skalaka U, Smailus DE, Schmerch A, Schein JE, Jones SJ and Marra MA.

Mammalian Gene Collection Program Team
Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clone distribution: MGC clone distribution information can be for through the I.M.A.G.B. Consortium/LLNL at: http://image.llnl.gov Series: IRBR Plate: 1 Row: h Column: 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue Procurement: Baylor Human Genome Sequencing Center CDNA Library Preparation: Baylor Human Genome Sequencing ocDNA Library Arrayed by: The I.M.A.G.E. Consortium (LIML) DNA Sequencing by: Baylor College of Medicine Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NIH-MGC Project URL: http://mgc.nci.nih.gov
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Submitted (29-APR-2004) National Institutes of Health, Mammalian
Gene Collection (MGC), Cancer Genomics Office, National Cancer
Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Director MGC Project.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Center code: BCM-HGSC
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                                                                     precursor"
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REFERENCE

AUTHORS

Hominidae; Homo.

1 (bases 1 to 531)

1 (bases 1 to 531)

Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RL, Feingold EA, Schuler GD, Altschul SF, Zeeberg Collins FS, Wagner L, Shenmen CM, Schuler GD, Altschul SF, Zeeberg B, Buetow KH, Schaefer CF, Bhat NK, Hopkins RF, Jordan H, Moore T, Max SI, Wang J, Hsieh F, Diatchenko L, Marusina K, Farmer AA, Rubi CM, Hong L, Stapleton M, Soares MB, Bonaldo MF, Casavant TL, Scheetz TE, Brownstein MJ, Usdin TB, Toshiyuki S, Carninci P, Prange C, Raha SS, Loquellano NA, Peters GJ, Abramson RD, Mullahy SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, Richards S, Worley KC, Hale S, Garcia AM, Gay LJ, Hulyk SW, Villalon DK, Muzny DM, Sodergren EJ, Lu X, Gibbs RA, Fahey J,

Zeeberg

SOURCE KEYWORDS

ORGANISM

Homo sapiens Homo sapiens (human) BC069523.1 GI:47479431

Eukaryota; Metazoa; Mammalia; Eutheria;

Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini;

VERSION ACCESSION

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                            Homo
IMAGE:7262093),
BC069523
                                           BC069523
                                                                                                    GAGGAGACAGAA 528
                                                                                                                                                                                                                                                                            GCCAATTCCTTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC
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               sapiens interleukin
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       US-10-789-129-26 (1-144)
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Center code: BCM-HGSC
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Contact: MGC help desk
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                                                                                                                                                                 Length:
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                                                                                                                                                                                                                                                                           Hominidae; Homo.

21 (bases 1 to 531)

22 Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Collins FS, Wagner L, Shenmen CM, Schuler GD, Altschul SF, Zeeberg B, Buetow KH, Schaefer CF, Bhat NK, Hopkins RF, Jordan H, Moore T, Max SI, Wang J, Heich F, Diatchenko L, Marusina K, Farmer AA, Rubin GM, Hong L, Stapleton M, Soares MB, Bonaldo MF, Casavant TL, Scheetz TE, Brownstein MJ, Usdin TB, Toshiyuki S, Carninci P, Prange C, Raha SS, Loquellano NA, Peters GJ, Abramson RD, Mullahy SJ, Bosak SA, McEwan DJ, McKernan KJ, Malek JA, Gunaratne PH, Richards S, Worley KC, Hale S, Garcia AM, Gay LJ, Hulyk SW, Villalon DK, Muzny DM, Sodergren EJ, Lu X, Gibbs RA, Fahey J, Helton E, Ketteman M, Maden A, Rodrigues S, Sanchez A, Whiting M, Madan A, Young AC, Shevchenko Y, Bouffard GG, Blakesley RW, Touchman JW, Green ED, Dickson MC, Rodriguez AC, Grimwood J, Schmutz J, Myers RM, Butterfield YS, Krzywinski MI, Skalska U, Smallus DE, Schnerch A, Schein JE, Jones SJ and Marra MA.

TM Mammalian Gene Collection Program Team Generation and initial analysis of more than 15,000 full-length human and mouse cDNA secuences
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Direct Submission
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Submitted (29-APR-2004) National Institutes of Health, Mammalian Submitted (29-APR-2004) National Institutes (29-APR-2004) Nati
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                                                                                                                              Director MGC Project.
                                                                                                                                                                                                                                                                    human and mouse cDNA sequences
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Baylor Human Genome Sequencing Center
cDNA Library Preparation: Baylor Human Genome Sequencing Center
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Baylor College of Medicine Human Genome
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Location/Qualifiers
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                                         AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp 60
                                                                                                                                                                                                  GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
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                                                                                                                                                                                                                                                                                                       CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
                                                                                                                         GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC
                                                                                                                                                                                                                                                                                    TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGAGATACGGGGCAGT
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/db_xref="MIM:605619"
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GAGGAGACAGAA
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Search completed: December 28, 2005, 15:08:47 Job time : 2947 secs

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Result
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-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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AAX77692
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Adm95042 Human Int
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ACF1325 ACD2535 ACF0040 ACA6043	B ACA66838 B ACA91232 B ACD81609 B ACF20413 B ACF20413 B ACF19799	6 ABS74446 6 ABL98214 6 ABL95703 7 ADY31957 8 ACA89679 8 ACA73689 8 ACA73689	8 ABV75154 12 ADL15289 6 ABQ74562 6 ABV73092 12 ADN12153 13 ADS31716 13 ADS92718	10 AAD61821 10 AD64668 11 AD646668 12 AD768066 13 ADR16327 13 ADR24325 13 ADR24325 14 AEA5608 14 AEA56986 14 AEA26786 14 AEA26786
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ALIGNMENTS

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ADM95042
ID ADM95042;
XX
AC ADM95042;
XX
C ADM95042;
XX

COLON cancer; interleukin 20 receptor alpha chain;
XX
COLON cancer; alpha chain; cytostatic; gene; ds; human.
XX

MIL-20 receptor alpha chain; cytostatic; gene; ds; human.
XX

FT CDS

FT CDS

AC ADM95042;
XX

FT CDS

AC ADM95042;

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In-2004

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tag SEQ

ID NO

428

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RESULT 2
ADV42800
ID ADV4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel agent for prevention and/or treatment of colon cancer, comprising of the interleukin 20 (II-20) receptor alpha chain, its partial peptide or salt. The invention may be useful for the development of compounds with a cytostatic activity. The invention is useful for preventing and/or treating colon cancer. A diagnostic agent is useful for diagnosis of colon cancer. By using II-20, II-20 receptor, its salt or its peptide, compounds that have altered binding property towards them can be efficiently screened. The present sequence is that of a gene which encodes the human Interleukin-20 protein and which is related to
                  ADV42800
                                               ADV42800 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 528 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Agent useful for prevention and/or treatment of colon cancer, comprises interleukin 20 (IL-20), IL-20 receptor alpha chain, IL-20 receptor beta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATTGGATTTTTCTGACATACGGGGCAGT
                                                                                                                                                                                                                                                              HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                           AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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                                                                                                                                             GluGluThrGlu
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                                                                                                                                                                                                                                                                                                         GCCAATTCCTTTCTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC
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                                               cDNA; 531
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a new microarray which comprises probes for genes involved in psychoneuroendocrinimmune (PNI) activity. The microarray is useful in diagnosing a condition associated with PNI activity, such as CFS, type-2 diabetes, allergic condition, inflammation, cancer and infection. The present sequence represents a psychoneuroendocrinimmune gene expressed sequence tag. Note the specificatio mentions SEQ ID NO of up to 3314 but only sequences up to SEQ ID NO 1829 are provided.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New microarray comprising probes for genes involved in psychoneuroendocrinimmune (PNI) activity, useful in diagnosing condition associated with PNI activity, e.g., inflammatory or i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 531 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microarray; psychoneuroendocrinimmune; chronic fatigue;
non-insulin dependent diabetes; allergy; immune disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 428;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-JUN-2003; 2003US-0475915P
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                                                                                              GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC
                                                                                                                                                                                  AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                                                                                                                             GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
                                                                                                                                                                                                                                                ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
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                   CC This invention describes a novel mammalian four alpha helix cytokine CC designated Zcytol0 which has vulnerary, cytostatic, anti-inflammatory, CC anti-asthmatic and haemostatic activity. Zcytol0 may be a growth and/or CC maintenance factor in the trachea and salivary glands, stomach, pancreas CC and muscle. Zcytol0 polynucleotides are useful as sources of primers and CC probes, and to determine if the Zcytol0 gene is present on chromosome 1, CC or to detect any mutations that may have occurred. Zcytol0 cc polynucleotides are also useful as gene therapy reagents. The Zcytol0 cc polynucleotides are also useful as gene therapy reagents. The Zcytol0 cc polypeptide, its fragments, and antibodies, as well as compounds cc identified as having binding affinity to Zcytol0, should be useful in the treatment of conditions associated with abnormal physiology or development, including abnormal proliferation, e.g. cancerous conditions, cc or degenerative conditions or altered immunity. The products are also cuscful for modulating cell proliferation, cell differentiation or cytokine production in the prevention or treatment of conditions or altered immunity. The products are also contains a provided by abnormal cell proliferation, cell differentiation or cytokine production.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New mammalian cytokine-like polynucleotide useful in the treatment asthma and bronchitis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inhibiting the growth and/or proliferation of cervical cancer cells, useful for treating cervical cancer or human papilloma virus infection, comprises bringing Interleukin-20 (IL-20) into contact with the cervical cancer cells.
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                                                                                             HisCysGlyGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                   AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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                                                                                                                       GCCAATTCCTTTCCTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC
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Human; cytokine-like polypeptide-10; Zcyto10; therapy; thrombocytopaenia; autoimmune disease; insulin-dependent diabetes mellitus; immune response; rheumatoid arthritis; multiple sclerosis; infection; radiation therapy; asthma; bronchitis; wound healing; periodontal disease; antiinflammatory; psoriasis; eczema; dry skin; oral cavity; muscle atrophy; haemostatic;
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25-NOV-1998;
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Novel human cytokine-like polypeptide-10, Zcyto10, useful for treating insulin-dependent diabetes mellitus, rheumatoid arthritis, multiple sclerosis, asthma, psoriasis and inhibiting cancer cell growth.

Disclosure; Col 29-30; Opp; English.

The invention relates to an isolated human cytokine-like polypeptide-10 (Zcytol0) polypeptide. Zcytol0 is useful for prevention or treatment of Cc conditions characterised by improper cell proliferation, cell crossis, etc., by inhibiting cellular cremmanced arthritis, multiple sclerosis, etc., by inhibiting cellular cremmanced arthritis, multiple sclerosis, etc., by inhibiting cellular cremmanced cremmanced in suseful for inhibiting cancer cell crossis, etc., by inhibiting cancer cell cremmanced in cancer patients who experience thrombocytophenia due to combat thus useful in cancer patients who experience thrombocytophenia due to cancer therapy or radiation therapy. Zcytol0 is useful in tracheacteric cremance or would repair of the tracheabronchial specific applications, such as in the cremance of debris or in treatment of applications, such as in the clearance of debris or in treatment of asthma, bronchits or other challing and promote regeneration of affected tissues which may be useful to the tracheabronchial tract. Zcytol0 may also enhance wound the treatment of periodontal disease. Zcytol0 can be useful to treat skin conditions such as psoriasis, eczema and dry skin in general. Zcytol0 is useful for regenerating gastrointestinal tract or þe

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                                                                                                cytokine-like polypeptide-10; Zcyto10; wound healing; platelet proliferation; wound repair; mucous production; asthma; bronchitis; tracheobronchial tract disease; periodontal disease; skin condition; psoriasis; eczema; dry skin; protein therapy; human; vulnerary; respiratory; antibronchitic; dental; dermatological; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel mammalian cytokine-like polypeptide-10 (Zcytol0) proteins and polynucleotides encoding such proteins. Sequences of the invention are useful for promoting the healing of wounds and for stimulating the proliferation of platelets. In particular Zcytol0 polypeptides may be used in trachea-specific or tracheobronchial-specific applications such as maintenance or wound repair of the tracheobronchial epithelium or cells underlying it; in regulating mucous production, mucocilary clearance of debris; in tracheobronchial tract. They are also used to promote regeneration of tracheobronchial tract. They are also used to promote regeneration of affected tissues which may be especially useful in the treatment of periodontal disease. Zcytol0 polypeptides can be used to treat skin conditions such as psoriasis, eczema and dry skin in general. They are also used in protein therapy. The present sequence is human Zcytol0 DNA.
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25-NOV-1998;
17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 926
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                                                                                                   GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC
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                                                                                             New ZcytolO polypeptides and polynucleotides, useful for treating autoimmune diseases (e.g. insulin dependent diabetes mellitus, multiple sclerosis or rheumatoid arthritis), or skin conditions (e.g. psoriasis)
                                                                                                                                                              WDI; 2004-552564/53.
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                                Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC (Zcyto10) proteins and their encoding polynucleotides. The invention is CC useful for treating conditions characterised by improper cell proliferation, cell differentiation or cytokine production, specifically CC proliferation, cell differentiation or cytokine production, specifically CC for treating autoimmune diseases such as insulin dependent diabetes cancer cell growth. The invention is also used to stimulate the immune CC cancer cell growth. The invention is also used to stimulate the immune CC tissues, to increase platelet count and to treat asthma and skin CC conditions such as psoriasis, eczema and dry skin. The invention acts as CC conditions such as psoriasis, eczema and dry skin. The invention acts as CC cytostatic, immunostimulant, vulnerary, antiasthmatic, antipsoriatic and CC dermatological agent. The invention is also used in gene therapy. The CC present sequence is human Zcytol0 longer form DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 926
                            Cytokine-like polypeptide-10; Zcyto10; therapy; asthma; infection; psoriasis; eczema; dry skin; wound healing; platelet proliferation;
                                                                          Human
                                                                                                         21-OCT-2004
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                                                                       cytokine-like polypeptide-10 (Zcyto10) long
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                                                                                                                                                                                                                                                                                                          GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet
                                                                                                                                                                                                                                                                                                                                                    CATTGTGGGGAGGAAGCAATGAAGAAATACAGCCAGATTCTGAGTCACTTTGAAAAAGCTG
                                                                                                                                                                                                                                                                                                                                                                    HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                                                                                                                GCCAATTCCTTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC
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25-NOV-1998;
15-APR-2003;
                                                                                                                                                                                                                                                                                                                               The invention relates to novel cytokine-like polypeptide-10 (Zcyto10) polypeptides and polynucleotides. Zcyto10 sequences are useful for treating sathma, microbial or viral infections, psoriasis, eczema or dry skin, for promoting the healing of wounds and for stimulating the proliferation of platelets. The present sequence is human Zcyto10 DNA.
                                                                                                                                                                                                                                                                                                              Sequence 926
                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 1; 32pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                 isolated antibody that binds to a ZcytolO polypeptide, useful ating asthma, microbial or viral infections, and for promoting ling of wounds, psoriasis, eczema or dry skin.
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DB; ADR16328, AE
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                                                                                                                                  ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
         AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
                                                 ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
                                                                                              AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                      GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
                                                                                                                                                                TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGACATACGGGGCAGT
                                                                                                                                                                            CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
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                                                                                                                                                                                                                                                                                                           The invention relates to a method of classifying a breast cancer patient according to prognosis by determining the similarity between the level of expression of each of five genes for which markers are listed in the specification, in a cell sample taken from the breast cancer patient, to control levels of expression for each respective five genes to obtain a patient similarity value. The methods are useful for classifying a breast cancer patient according to prognosis. Kits and computer program products are useful for data analysis using the diagnostic, prognostic and statistical methods of the invention. This sequence corresponds to a marker used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Classifying a breast cancer patient according to prognosis comprises determining the similarity between the level of expression of each of five genes in a cell sample taken from patient, to control levels.
                                                                                                                                                                                                                                                               Sequence
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25-NOV-1998;
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GluGluThrGlu

144 572

RESULT 11
AEA50048
ID AEA50
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AC AEA50
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DT 11-AU
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DE Full

standard;

CDNA;

ВP

AEA50048 sta AEA50048; 11-AUG-2005 Full length

IL-20 coding sequence

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Percent Similarity:
Best Local Similarity:
Query Match:
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P-PSDB;
                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a mammalian cytokine-like polypeptide, Zcytol0 and its encoding polymucleotide. The invention is useful for modulating cell proliferation, cell differentiation or cytokine production in the prevention or treatment of conditions such as autoimmune diseases (e.g. diabetes, multiple sclerosis, rheumatoid arthritis, asthma or psoriasis) and cancer or wounds. The invention is also useful in gene therapy. The present sequence is the human Zcytol0 protein encoding DNA. This Zcytol0 gene is located on chromosome 1q32.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antibody that selectively binds to a ZcytolO polypeptide useful modulating cell proliferation, cell differentiation or cytokine production in the prevention or treatment of e.g. autoimmune disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure;
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DB; ADS18363, ADS18373.
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                 ArgvalPheLy8AsnTyrGlnThrProAspHisTyrThrLeuArgLy8IleSerSerLeu
                                                                GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
  AGGGTATTTAAAAACTACCAGACCCCTGACCATTATACTCTCCGGAAGATCAGCAGCCTC
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Matches:
Conservative:
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
                                                                                                                                           This sequence encodes full length interleukin-20 (IL-20), longer form IL-20 (20x1. The method of the invention for producing an antibody to IL-20 (comprises inoculating an animal with an IL-20 polypeptide which elicits an immune response in the animal. The produce the antibody, and isolating the antibody from the animal. The antibody is useful for treating a compatibility produced the antibody from the animal. The antibody is useful for treating a compatibility produced with IL-20 activity such as compatible of the antibody is useful for treating a compatibility produced with IL-20 activity such compatible of the antibody is useful for treating a manufactory bowel compatible of IL-20 is useful for treating a manufaction such as contact and psoriasis, or acute inflammatory condition such as contact antibody and infectious disease. An compatible of IL-20 is useful for treating a manufact with an compatible of IL-20 is useful for treating a manufact with an confident of IL-20 is useful for treating a manufact that the compatible or polypeptide fragment of IL-20 and where the inflammatory activity of IL-20 antibody. The antibody or antibody fragment, produced using the above contactivity of IL-20. IL-20 and inhibits or reduced the pro-inflammatory activity of IL-20 and inhibits or reduced the pro-inflammatory contactivity of IL-20. IL-20 induced proliferation or differentiation of inhibited mains the antibody antibody can inhibit or reduced the pro-inflammatory antibody inhibited mains the antibody and inhibits or reduced the pro-inflammatory antibody antibody antibody antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Producing an antibody to an interleukin-20 (IL-20) polypeptide, useful for treating inflammatory diseases e.g., Crohn's disease, by inoculating an animal with the polypeptide eliciting an immune response to produce
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antiarthritic; Dermatological; Antipsoriatic; Antibacterial; Immunosuppressive; Antimicrobial; II-Antagonist-20; interleukin-20; II-20; inflammation; inflammatory bowel disease; ulcerative colitis; Crohns disease; arthritis; atopic dermatitis; psoriasis; endotoxemia; septicemia; toxic shock syndrome; infectious disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; SEQ ID NO 1; 268pp; English.
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24-MAR-2004;
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Antiarthritic;
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                                                                                                                                                                                                                                                                                                                                                                                           The invention provides a new purified polynucleotide that comprises the concleic acid sequence A-B-C-D-E; where: A- start codon, B- polynucleotide concleic acid sequence of 13 nucleic acids, where the sequence has homology with 13 concleic pair (bp) sequence, where at least 5 nucleic acids are identical to code for a stop codon, C- polynucleotide sequence comprising 10, 13, 16 concleic acids of a 19 bp sequence (ABY75139), and where the nucleic acid sequence does not code for a stop codon, C- polynucleotide sequence comprising 10, 13, 16 concleic acids of a 19 bp sequence (ABY75140), D- polynucleotide concleic acids of a 19 bp sequence (ABY75140), D- polynucleotide concleic acid from 9 sequences (ABY75144-150) where at least 50% nucleotides are either adenine or thymine, and where only E encodes a stop codon. The colynucleotide is useful for producing a nucleic acid construct for copyracsion of an amino acid sequence of interest. The present sequence corpresents a human interleukin-20 (IL-20) codon optimised fragment for construct for expression in E. coli
                                                                                                                                                                                                                          US-10-789-129-26 (1-144) x ABV75151 (1-462)
                                                                                                                                                                                                                                                                                  Best Local Similarity:
                                                                                                                                                                                                                                                                                               Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polynucleotide, useful for producing a nucleic acid construct expression of an amino acid sequence of interest.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein synthesis; data mining; recombinant; interleukin-20; IL-20;
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                                                                                                                                                                                                                                                                                                                                                                   Sequence 462
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                                                                                                                                                                  The invention relates to Syn 1 and Syn 2 tight regulated synthetic promoters and methods for enhancing the expression of recombinant proteins. The Syn 1 and Syn 2 tight regulated synthetic promoters are useful for controlling uninduced, leaky expression of proteins that are toxic and interfere with production of recombinant protein from an Escherichia coli expression system. They are useful for controlling basal expression of target genes. The present sequence is the human interleukin [II] 20 DNA. This sequence is used in the comparison of promoter
                                                                                                   Sequence 462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New Syn 1 and Syn 2 tight regulated synthetic promoters, useful for controlling uninduced, leaky expression of proteins that are toxic and interfere with production of recombinant protein from an Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; SEQ ID NO 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression
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DB; AEB26650.
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/product= "Human interleukin
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                                          New polynucleotide, useful for producing a nucleic acid expression of an amino acid sequence of interest.
                                                                                                                                                              Pownder TA,
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Example 1; Page 37; 40pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention provides a new purified polynucleotide that comprises the nucleic acid sequence A-B-C-D-E; where: A= start codon, B= polynucleotide sequence of 13 nucleic acids, where the sequence has homology with 13 base pair (bp) sequence, where at least 5 nucleic acids are identical to a 13 bp sequence (ABV75139), and where the nucleic acid sequence does not code for a stop codon, C= polynucleotide sequence comprising 10, 13, 16 or 19 nucleic acids of a 19 bp sequence (ABV75140), D= polynucleotide sequence comprising 8 bp (ABV75141), E= polynucleotide sequence selected from 9 sequences (ABV75141-150) where at least 50% nucleotides are either adenine or thymine, and where only E encodes a stop codon. The polynucleotide is useful for producing a nucleic acid construct for expression of an amino acid sequence of interest. The present sequence represents a sequence comprising an expression cassette (EC) polynucleotide sequence inserted upstream of a human interleukin-20 (IL-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 513 BP; 143 A; 145 C;
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                                                                                         GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
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                                          GluGluThrGlu 144
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-Q=/Ggn2 1/USPTO spool p/US10789129/runat 23122005 152720 20276/app query.fasta_1.327
-DB=EST -QFWT=fastap - SUFFIX.p201.rst - MINMATCH=0.1 - LOOPEXT=0
-UNITS=bits -START1 - LEND=-1 -MAYRIXEblusde - THANS=human40.cdi - LIST=45
-DCCALIGN=200 -THR SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFWT=pto -NORM-ext -HEAPSIZE=500 -MINLENS - NAXLEN-200000000
-USER=US10789129 GCGN_1 1 8010 garunat 23122005 152720 20276 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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ALIGNMENTS

REFERENCE AUTHORS RESULT 1 AY402693 LOCUS REFERENCE AUTHORS SOURCE ORGANISM VERSION KEYWORDS COMMENT DEFINITION ACCESSION JOURNAL PUBMED TITLE JOURNAL TITLE Submitted (16-NOV-2003) Celera Genomics, 45 West Gude Drive, Rockville, MD 20850, USA
This sequence was made by sequencing genomic exons and ordering 2 (bases 1 to 531)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, J. Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
Direct Submission gene trios Science 302 (5652), 1960-1963 Hominidae; Homo.

1 (bases 1 to 531)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Clark, A.G., Glanowski, S., Nielson, R., Lu, F., Murphy, B., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
Adams, M.D. and Cargill, M.
Inferring nonneutral evolution from human-chimp-mouse orthologous Homo sapiens (human) genomic survey sequence.
AY402693 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; AY402693.1 GI:39758676 GSS. Homo sapiens IL20 gene, AY402693 14671302 531 bp VIRTUAL (2003) TRANSCRIPT, partial sequence, DNA linear Kejariwal, A., GSS 15-DEC-2003

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RESULT 2
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         Hominidae; Pan.

(bases 1 to 531)

(bases 1 to 531)

Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A.,

Clark, A.G., Glanowski, S., Nielson, R., Lu, F., Murphy, B.,

Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B.,

Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J.,

Adams, M.D. and Cargill, M.

Inferring nonneutral evolution from human-chimp-mouse orthologous
                                                                                                                                                                 GSS
                                                                                                                                                                            Pan troglodytes IL20 gene, genomic survey sequence. AY402694 AY402694.1 GI:39758677
gene trios
                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
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Pan troglodytes
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/mo1_type="genomic DNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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/locus_tag="HCM1309"
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99.47%
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PUBMED
REFERENCE
AUTHORS
ACCESSION
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LOCUS
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Best Local Similarity:
Query Match:
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genomic survey sequence. AY402695
AY402695.1 GI:39758678
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Mus musculus (house mouse
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14671302
2 (bases 1 to 531)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Trodd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
                                                                    AY402695 531 bp
Mus musculus IL20 gene, VIRTUAL
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                                                                                                                                                                                            GAACCTCAGGCAGCAGTTGTGAAGGCTTTGGGGGGAACTAGACATTCTTCTGCAATGGATG
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/gene="IL20"
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/mol_type="genomic DNA"
/db_xref="taxon:9598"
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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/db_xref="taxon:10090"
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                      Submitted (16-APR-2002) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokoham Kanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/, Tel:81-45-503-9222, Fax:81-45-503-9216)
                                                                                                                                                                                                                             Adachi, J., Aizawa, K., Akimura, T., Arakawa, T., Bono, H., Carninci, P. Pukuda, S., Furuno, M., Hanagaki, T., Hara, A., Hashizume, W., Hayashida, K., Hayatsu, N., Hiramoto, K., Hiramoka, T., Hirozane, T., Harai, F., Imotani, K., Ishii, Y., Itoh, M., Kagawa, I., Kasukawa, T., Katoh, H., Kawai, J., Kojima, Y., Kondo, S., Konno, H., Kouda, M., Koya, S., Kurihara, C., Matsuyama, T., Miyazaki, A., Murata, M., Nakamura, M., Nishi, K., Nomura, K., Numazaki, R., Ohno, M., Ohsato, N., Okazaki, Y., Saito, R., Saitoh, H., Sakai, C., Sakai, K., Sakazume, N., Sano, H., Sasaki, D., Shibata, K., Shinagawa, A., Shiraki, T., Sogabe, Y., Tagama, M., Tagawa, A., Takahashi, F., Takaku-Akahira, S., Takada, Y., Tanaka, T., Tomaru, A., Toya, T., Yasunishi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Group Phase I & II Team.
Analysis of the mouse transcriptome of 60,770 full-length cDNAs
Nature 420, 563-573 (2002)
6 (bases 1 to 1189)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The FANTOM Consortium and the RIKEN Group Phase I & II Team.
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Mus musculus adult male eyeball cDNA, RIKEN full-length enriched
library, clone:7530415H19 product:interleukin 20, full insert
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High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
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Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         URL:http://genome.gsc.riken.jp/
URL:http://fantom.gsc.riken.jp/.
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Please visit our web site for further details.
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/clone_Tib="RIKEN full-length enriched mouse cDNA library"
/dev_stage="adult"
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/db_xref="FANTOM_DB:7530415H19"
/db_xref="taxon:10090"
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/mol_type="mRNA"
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Contact: Yoshihide Hayashizaki
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Konno, H., Fukunishi, Y., Shibata, K., Itoh, M., Carninci, P., Sugahara, Y. and Hayashizaki, Y. Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Kondo, S., Shinagawa, A., Saito, T., Kiyosawa, H., Yamanaka, I., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K. and Hayashizaki, Y.
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Arakawa, T., Carninci, P., Fukuda, S., Furuno, M., Hanagaki, T.,
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The Institute Oppsical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045,
Tel: 81-45-503-9222
Fax: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences, Mamm. Genome. 12, 673-677 (2001)
Please visit our web site (http://genome.gsc.riken.go.jp) for
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RIKEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome 10 (11), 1757-1711 (2000)
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/note="Site 1: Sall; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was
                                                                                                                                                                                                                   /tissue type="eyeball"
/dev_stage="adult"
/lab_host="DH10B"
/clone lib="RIKEN full-length enriched, adult male
eyeball"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="mRNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                mRNA seq
CR985515
                                                                                        Human T-Lymphocytes library
Unpublished (2005)
Contact: Inge Arlart
                                              RZPD Deutsches Ressourcenzentrum fuer Genomforschung
Heubnerweg 6, D-14059 Berlin, Germany
Email: www.rzpd.de
RZPDLIB; (Human T-Lymphocytes) RZPD LIB No.9016
http://www.rzpd.de/cgi-bin/products/set.cgi?libNo=9016 Contact:
                                                                                                                                                                                         Hominidae; Homo
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                                                                                                                                                                     (bases 1 to 647)
                                                                                                                                            Korn, B.
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                                RZPDp9016M1629.
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RZPD Deutsches Ressourcenzentrum fuer Genomforschung
Heubnerweg 6, D-14059 Berlin, Germany
Tel: +49 30 32639 100
Fax: +49 30 32639 111
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 GluGlu 142
                                            GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
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                            GAGGTCCACGCTGCCATTAAATCCCTGGGAGAGCTCGACGTCTTTCTAGCCTGGATT
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/lab_host="DH10B"
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Hara, A., Hiramoto, K., Hori, F., Ishli, Y., Ito, M., Kawai, J.,
Konno, H., Kouda, M., Koya, S., Mateuyama, T., Miyazaki, A., Nomura, K.,
Ohno, M., Okazaki, Y., Okido, T., Saito, R., Sakai, C., Sakai, K.,
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Takeda, Y., Tanaka, T., Toya, T., Muramatsu, M. and Hayashizaki, Y.
RIKEN Mouse ESTS (Arakawa, T., et al. 2001)
On Jun 14, 2000 this sequence version replaced gi:8522215.
Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sugahara, Y. and Hayashizaki, Y. Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Kondo, S., Shinagawa, A., Saito, T., Kiyosawa, H., Yamanaka, I., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K. Hayashizaki, Y.
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Carninci, P., Shibata, Y., Hayatsu, N., Sugahara, Y., Shibata, K.,
Itoh, M., Konno, H., Okazaki, Y., Muramatsu, M. and Hayashizaki, Y.
Normalization and subtraction of cap-trapper-selected cDNAs to
prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. . 10 (10), 1617-1630 (2000)
wagi, K., Pujiwake, S., Inoue, K., Togawa, Y., Izawa, M., Ohara, B.,
watahiki, M., Yoneda, Y., Ishikawa, T., Ozawa, K., Tanaka, T.,
Watauura, S., Kawai, J., Okazaki, Y., Muramatsu, M., Inoue, Y., Kira, A.
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The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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EST.
Mus musculus (house mouse)
Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences Mamm. Genome. 12, 673-677 (2001)
Please visit our web site (http://genome.gsc.riken.go.jp/) for further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                          prepare mouse tissues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and Hayashizaki,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory for Genome Exploration Research Group, RIKEN Genomic
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81-45-503-9216
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                                         /dev_stage="adult"
/lab_host="DH10B"
                                                                                             /tissue_type="eyeball"
                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
'clone_lib="RIKEN full-length enriched, adult male
                                                                                                                                          sex="male"
                                                                                                                                                                                       clone="7530415H19"
                                                                                                                                                                                                                                               strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
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                                                                                                                                                                                                                 _xref="taxon:10090"
                                                                                                                                                                                                                                                                                        _type="mRNA"
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/note="Site 1: SalI; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was primed with a primer [5]
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RESULT 8 CR992594 밁 Percent Similarity: Best Local Similarity: US-10-789-129-26 (1-144) x BB064801 (1-695) Query Match: Pred. Alignment Scores: 242 AGATGGATGGAGGAG 256 138 GlnTrpMetGluGlu 142 182 ATAGAGTTGGAACTTCAGGCÁGCGGTGGTAAAGGCTTTGGGAGAACTAGGCÁTTCTTCTG 118 GluLysLeuGluProGlnAlaAlaValLysAlaLeuGlyGluLeuAspIleLeuLeu 123 ATGGCATGTCA-TGTGGGGAAGAAGCAATGGAGAAATACAACCAAATTCTGAGTCACTTC 98 MetThrCysHisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPhe 63 78 SerSerLeuAlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHis 58 TyrLeuAspArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIle w CR992594 RZPD no.9016 Homo sapiens cDNA clone RZPDp9016C1545 5', AGCAGCCTCGCCAAATCCTTTCTTATCATCAAGAAGGACCTCTCAGTCTGTCATTCTCAC TATCTGGACAGGGTATTCAAAGTCTACCAGACCCCTGACCNCCATACCCTGAGAAAGATC 6.17e-30 328.00 38.24% 80.00% 43.56% Mismatches: Indels: Gaps: Length: Matches: Conservative: 117 241 181 122 97 62

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VERSION KEYWORDS

ACCESSION DEFINITION

mRNA sequence. CR992594

CR992594.1

GI:68286479

SOURCE

ORGANISM

Homo sapiens

Homo sapiens (human)

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;

JOURNAL COMMENT

Unpublished (2005

and Korn, B.

(bases 1 to 699) il,O., Ebert,L., Hennig,S., T-Lymphocytes library

Schneider, D

Contact: Inge Arlart

RZPD Deutsches Ressourcenzentrum fuer Genomforschung GmbH Heubnerweg 6, D-14059 Berlin, Germany Email: www.rzpd.de

TITLE AUTHORS

REFERENCE

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-789-129-26 (1-144) x CR992594 (1-699)
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Tel: +49 30 32639 110
Fax: +49 30 32639 111
www.rzpd.de
This clone is available
                                                                                                                                                                                                                                                                                                                                                                                                                             21 ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This clone is available from RZPD;

http://www.rzpd.de/cgi- bin/products/cl.cgi?CloneID=RZPDp9016C1545

contact RZPD (product- support@rzpd.de) for further information.

Primer name: qe3 4 , Primer sequence: CGGATAACAATTCACACAG.

Location/Qualifiers
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RZPD
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http://www.rzpd.de/cgi-bin/products/set.cgi?libNo=9016 Contact:
                                                                                                       HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                               ATCATTAAGCCCTTAGATGTGTGCGTGACCAAGAACCTCCTGGCGTTCTACGTGGAC
                      GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
                                                                                                                                                                                                                                                                                                                                                                                           ATCCAAGCTAAGGACACCTTCCCAAATGTCACTATCCTGTCCACATTGGAGACTCTGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
                                                                                                                                                      GCCAACTCTTTCCTCTACATGCAGAAAACTCTGCGGCAATGTCAGGAACAGAGGCAGTGT
                                                                                                                                                                               AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
                                                                                                                                                                                                                                                          ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
GAGGTCCACGCTGCCATTAAATCCCTGGGAGAGCTCGACGTCTTTCTAGCCTGGATT
                                                                           CACTGCAGGCAGGAAGCCACCAATGCCACCAGAGTCATCCATGACAACTATGATCAGCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="RZPDp9016C1545"
/tissue_type="T-Lymphocytes"
/dev_stage="adult"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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43.56%
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Matches:
Conservative:
Mismatches:
Indels:
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DEFINITION
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KEYWORDS
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Tissue Procurement: Dr. Gary W. Hunninghake, U of I
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           762 bp mRNA linear EST 05-AUG-
UI-H-FT1-bic-n-07-0-UI.81 NCI CGAP FT1 Homo sapiens cDNA clone
UI-H-FT1-bic-n-07-0-UI 3', mRNA sequence.
CA309757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Gene Index
Unpublished (1997)
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1 (bases 1 to 762)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST
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POLYA=Yes.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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           /clone="berlow white recursory conditions (blasmacia) with a modified polylinker; Site_1: Ecc I; Site_2: Not I; No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_cype= ".....9606"
/db_xref="taxon:9606"
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/lab_host="DH10B (Life Technologies)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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TISSUE=Human
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PUBMED
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23
                                                                                                                                                                                                                                                                                1 (bases 1 to 564)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W.Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Simpson,A.J.
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM4&t2=PM4-KT0045-
151200-006-g02&t3=2000-12-15&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PM4-KT0045-151200-006-g02
BF738279
                                                                                                            Brazil
Tel: +55-11-2704922
                                                                                                                                                            Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                   Shotgun sequencing of the human transcriptome with ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BF738279.1 GI:12065059
EST.
                                                                                                                                               Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                                                                                                                                                                                   sequence tags
                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                               Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GlnAlaAlaValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMetGluGlu 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCysHisCys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGGCAGGAAGCCACCAATGCCACCAGAGTCATCCATGACAACTATGATCAGCTGGAGGTC 463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeuGluPro 122
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                                                                                            +55-11-2707001
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TAG_SEQ=GGCCATGCCG"
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275.00
65.00%
45.00%
36.52%
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KT0045 Homo
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Matches:
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source

/organism="Homo sapiens" /mol_type="mRNA" /db_xref="taxon:9606"

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REFERENCE
AUTHORS
TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       509 CAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGACAGGGTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                43 LysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAspArgVal 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CA306526
UI-H-FT1-bhw-m-10-0-UI.sl NCI CGAP_FT1 Homo sapiens
UI-H-FT1-bhw-m-10-0-UI 3', mRNA sequence.
                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Gary W. Hunninghake, U of I
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be obtained
from Dr. M. Bento Soares, bento-soares@uiowa.edu
                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Primates; Cata
                                                                                                                                                                                                                     Unpublished (1997)
                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                   Hominidae; Homo.
1 (bases 1 to 748)
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                                                                                                                                                                                                  Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
                                     Seq primer: M13 FORWARD
                  POLYA=Yes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TCCTTTCTTACCATCAAGAAGGACCTCCGGCTCTGT
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                                                                                                                                                                                                                                        Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Torgan: bladder tumor; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-Tibrary was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
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Location/Qualifiers
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36.25%
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Matches:
Conservative:
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Indels:
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                                                                                                                                                                                                                                                                                                                                                 Catarrhini;
                                                                                                                                                                                                                                                                 Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                  Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cDNA clone
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US-10-789-129-26 (1-144) x CA306526 (1-748)
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Best Local Similarity:
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                                                                                                                          624
                                                                                                                                                                       684
                                                                                                                                                                                                                  744 AATGTCACTATCCTGTCCACATTGGAGACTCTGCAGATCATTAAGCCCTTAGATGTGTGC
                                                                                                                                                69
                                                                                                                                                                                              49
AlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMetGluGlu 142
                                  GCCACCAGAGTCATCCATGACAACTATGATCAGCTGGAGGTCCACGCTGCTGCCATTAAA
                                                                                                                         LysAspLeuArgLeuCysHisAlaHisMetThrCysHisCysGlyGluGluAlaMetLys 108
                                                      LysTyrSerGlnIleLeuSerHisPheGluLysLeuGluProGlnAlaAlaValValLys 128
                                                                             TGCGTGACCAAGAACCTCCTGGCGTTCTACGTGGACAGGGTGTCAAAGGATCATCAGGAG
                                                                                                                                                                                          CysLeuLeuArgHisLeuLeuArgLeuTyrLeuAspArgValPheLysAsnTyrGlnThr 68
                                                                                                                                                                                                                                        AspIleArgIleLeuArgArgThrGluSerLeuGlnAspThrLysProAlaAsnArgCys
                                                                                                                                                                                                                                                                                                                                                                                                                     TAG_TISSUE=Human Lung Alveolar Macrophage
TAG_LIB=UI-H-FT1
TAG_SEQ=GGCCATGCCG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev_stage="Adult"
/lab_host="DH10B (Life Technologies)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="UI-H-FT1-bhw-m-10-0-UI"
/tissue_type="Alveolar Macrophage"
                                                                                                                                                                                                                                                                                                4.69e-22
265.00
65.79%
44.74%
35.19%
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Matches:
Conservative:
                                  449
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Query

No.:

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JOURNAL
COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/Carninci.p., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y. Normalization and subtraction of cap-traper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes. Genome Res. 10 (10), 1617-1630 (2000) wagi,K., Fujiwake,S., Inoue,K., Togawa,Y., Izawa,M., Ohara,E., Watshiki,M., Yoneda,Y., Ishikawa,T., Ozawa,K., Tanaka,T., Matsuura,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A., Prepri incorrects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory for Genome Exploration Research Group, RIKEN Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-Tel: 81-45-503-9222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hara,A., Hiramoto,K., Hori,F., Ishii,Y., Ito,M., Kawai,J., Konno,H., Kouda,M., Koya,S., Matsuyama,T., Miyazaki,A., Nomura,K., Ohno,M., Okazaki,Y., Okido,T., Saito,R., Sakai,C., Sakai,K., Sano,H., Sasaki,D., Shibata,K., Shinagawa,A., Shiraki,T., Sogabe,Y., Suzuki,H., Tagami,M., Tagawa,A., Takahashi,F., Takeda,Y., Tanaka,T., Toya,T., Muramatsu,M. and Hayashizaki,Y. RIKEN Mouse ESTs (Arakawa,T., et al. 2001)

Contact: Yoshihide Hayashizaki
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Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences. Mamm. Genome. 12, 673-677 (2001)
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Computer-based methods for the mouse full-length cDNA
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e mouse tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               encyclopedia: real-time sequence clustering for construction of enonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Kondo, S., Shinagawa, A., Salto, T., Kiyosawa, H., Yamanaka, I., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RIKEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome 10 (11), 1757-1771 (2000)
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
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BB639581.1 GI:16474967
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/note="Site 1: Sall; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory
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RIKEN full-length enriched, 7 days neona
cDNA clone A730059E15 5', mRNA sequence.
                                                                                                                                                                                               /tissue_type="cerebellum"
/dev_stage="7 days neonate"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                               organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                      clone="A730059E15"
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                                                                                                                                                                 lib="RIKEN full-length enriched, 7 days neonate
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TITLE
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Query Match:
                                                                                                       COMMENT
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                                                                                                                                                                                                                                                                                                                                AGENCOURT 18791077 NICHD_XGC_Te2N Xenopus laevis cDNA clone IMAGE.7203819 5', mRNA sequence.
                National Cancer Institute / NIH
Bldg. 31 Rm10A07 Bethesda, MD 20892
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Igor B. Dawid
                                                                                                                    NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection
Unpublished (1999)
                                                                                    Contact: Daniela S. Gerhard, Ph.D. Office of Cancer Genomics
                                                                                                                                                                                                                                                                                                    CK800046.1 GI:42812042
                                                                                                                                                                           Xenopodinae; Xenopus; Xenopus.
1 (bases 1 to 897)
                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
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Xenopus laevis
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cDNA Library Preparation: Express Genomics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTCTTAAGGAGAATCAGCAGCATTGCCAACTCTTTCCTCTGCGTGCAGAAATCTCTCTGGAG 212
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AGGACTCTGCTGGAGCAGGAGACACCAGGAGAGGCCTGGAGATGTGTGCTGCATGACCAAC 92
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                                                                                                                          141 GluGlu 142
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                                                                                                                                                                                                                                                                                               101 HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                                           531 GCTAACTCTGTCCTGGGTCTAAAGATAGAGCTCAAGCATTGTCATTCTTCTATGAGGTGT
                                                                                                                                                                                                                                                                                                                                                                                                                            471 AATATTTTCCCAAAAACTGGAAATCTCTTCAATTAAGGAGCAGAAAGGTCTTAATCATTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
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DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM15082 row: e column: 01
                                                                                                                                                                    GAGACAGAAGCAGCTATTATTAAAGCTATTGGAGACTTGAATATCCTGATTCGCTGGTTG
                                                                                                                                                                                                                                                        CCCTGTGGTGATCAGTCACACAAAATTATGGAAGACTTTAGGGAAACCTTTTACCAGATG
                                                                                                                                                                                                                                                                                                                                                                                AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                was primed using oligo-dT primer:
5'-pGACTAGTTCTAGATCGCGGGCGGCCC(T)25-3' and cloned into
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/lab host="DHIOB TonA"
/clome_lib="NICHD XGC_Te2N"
/clome_lib="NICHD XGC_Te2N"
/note="Organ: testis; Vector: pExpress-1; Site_1: EcoRV;
Site_2: NotI; RNA obtained from 6 adult male testis. cDNA
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Best Local Similarity:
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      US-10-789-129-26 (1-144) x CD366952 (1-658)
                                                                                                                                                                                                                                        Alignment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Gary W. Hunninghake, U of I
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Distribution information can be found at http://genome.uiowa.edu/distribution/cgap.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hominidae; Homo.

1 (bases 1 to 658)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD366952.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tumor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer: M13 FORWARD
                                                                                                                                                                                                                                                                                                                                                                                                                                     the conditions, other donor macrophages in different conditions). The mRNA samples were pooled for library construction. Control 0 hours; control 3 hours; control 24 hours; LPS 100 ng/ml, 3 hours; LPS 100 ng/ml, 24 hours; PMA 10 ng/ml, 3 hours; PMA 10 ng/ml, 24 hours; Klebsiella moi 10, 3 hours; PMA 10 ng/ml, 24 hours; Staph aureus moi 10, 3 hours; Staph aureus moi 10, 24 hours; Adenoviral vector (Ad5 CMV eGFP), moi 500, 3 hours; wt adenovirus moi 500, 3 hours; wt adenovirus moi 500, 3 hours; wt adenovirus to 500, 24 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 3 hours; must adenovirus + LPS 3 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 24 hours; wt adenovirus + LPS 26 hours; wt adenovirus + LPS 27 hours; wt adenovirus + LPS 28 hours; wt adenovirus + LPS 29 hours; wt adenovirus + LPS 29 hours; wt adenovirus + LPS 20 hours; wt adenovirus + LPS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site 1: EcoR I; Site 2: Not I; NCI_CGAP_FTZ is a subtracted cDNA library constructed from a pool of 81 RNA samples from Alveolar Macrophages challenged with different treatments. The mRNA samples
                                                                                                                                                                                                                                                                                                                         TAG_TISSUE=Human Lung
TAG_LIB=UI-H-FT2
TAG_SEQ=GGCCATGCCG*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone lib="NCI_CGAP_FT2"
/note="Organ: Lung; Vector: pT7T3-Pac
modified polylinker; Site 1: EcoR_I; S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  were a mixture of these conditions (times refer to incubations following isolation by bronchoalveolar lavage)
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/db_xref="taxon:9606"
/clone="UI-H-FT2-bjp-i-22-0-UI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="Alveolar Macrophage"
/dev_stage="Adult"
/lab_host="DH10B (Life Technologie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (some normal donor macrophages were cultured in some of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: black009@niehs.nih.gov
Clone is available through Re
Parkway, Huntsville, AL 35901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Perry J. Blackshear
Office of Clinical Research and Laboratory of Signal Transduction
National Institute of Environmental Health Sciences
A2-05 NIEHS, 101 Alexander Drive, Research Triangle Park, NC 27709,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Blackshear, P.J., Lai, W.S., Thorn, J.M., Kennington, E.A., Staffa, N. Jr., Moore, D.T., Bouffard, G.G., Beckstrom-Sternberg, S.M., Touchman, J.W., Bonaldo, M.F. and Forers, M.B.
The NIEHS Xenopus maternal EST project: interim analysis of the first 13,879 ESTs from unfertilized eggs
Gene 267 (1), 71-87 (2001)
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bl54f02.w1 Blackshear/Soares nor
laevis cDNA clone PBX0054F02 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phone 800-533-4363 ext.cdna, fax 256-536-9016 att:cdna, email cdna@resgen.com
                                                                                                                                                                                                                                                                                                                                                                                                                      FORWARD: TGTAAAACGACGGCCAGT
BACKWARD: CAGGAAACAGCTATGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR PRimers
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Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 919 541-4899
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1 (bases 1 to 538)
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      /veri Type="unfertilized egg"
/dev stage="unfertilized egg"
/lab_host="DH10B"
/clone lit
                                                                                                                                                                                                      /organism="Xenopus laevis"
/mol_type="mRNA"
/db_xref="taxon:8355"
                                                                                                                                                 'sex="female"
                                                                                                                                                                                   clone="PBX0054F02"
                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                         CAGGAAACAGCTATGACC
clone_lib="Blackshear/Soares normalized Kenopus
                                                                                                                                                                                                                                                                                                                                                                                          row: F column:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mesobatrachia; Pipoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bp mRNA linear
normalized Xenopus egg
5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            performed by National Center (NISC).
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library Xenopus

EST 26-APR-2001

418

Euteleostomi;

Pipidae;

Staffa, N.G.

Institutes

Memorial

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
Search completed: December 28, 2005, 16:12:12 Job time : 3796 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-789-129-26 (1-144) x AW637118 (1-538)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
                                                                                         322
                                                                                                                                                                                                                                                                        262
                                                                                                                                                                                                                                                                                                                                                                                                                                                        142
                                                                                                                                                                                                                         104 GluGluAlaMetLysLysTyrSerGlnIleLeuSerHispheGluLysLeuGluProGln 123
                                                                                                                                                                                                                                                                                                                                                               202 CCAAAACTGGAAATCTCTTCAATTAAGGAGCAGAAAGGTCTTAATCATTTÄGCTÄÄCTCT 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44
                                                                                                                                                                                                                                                                                                  84 PheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCysHisCysGly 103
                                                                                                                                                                                                                                                                                                                                                                                                         64 LysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeuAlaAsnSer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24 LysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGlnAspThrLys 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GATCAGTCACAAAATTATGGAAGACTTTAGGGAAACCTTTTACCAGATGGAGACAGAA 381
                                                                                                                                                                                                                                                                        GTCCTGGGTCTAAAGATAGAGCTCAAGCATTGTCATTCTATGAGGTGTCCCTGTGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAGGATGTGATTACAGATGTCAGCCTCCTG---AAAGCCAAAGTGCTGAATCAGATACAT 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="vector: pT7T3-Pac; Site_1: EcoRI; Site_2: NotI; PolyA-selected mRNA was prepared from unfertilized Xenopus laevis eggs. The library was constructed in the vector pT7T3-Pac as described in Bonaldo, M.F., Lennon, G. and Soares, M.B. 'Normalization and subtraction: two approaches to facilitate gene discovery', Genome Research 6:791-806, 1996. The first strand synthesis used a NotI-dT18 primer; double stranded cDNAs were ligated to EcoRI adapters, digested with NotI, and directionally cloned into the NotI and EcoRI-digested pT7T3-Pac vector. The library contained approximately 7.2 X 10°5 recombinants, with average insert sizes of 1-1.5 kb."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             library"
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206.50
59.66%
32.77%
27.42%
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Matches:
Conservative:
Mismatches:
Indels:
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39
47
1
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Result
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-Q=-/cgn2 1/USPTO spool p/US10789129/runat 23122005 152721 20318/app query.fasta_1.327
-DBs-Published Applications NA Main -OPMT=fastap -SUPFIX=p2n.rnpbm
-MINMATCH=0.1 -LOOPCLAT - COOPEXT=0 - UNITS=blsts -START=1 -END=-1
-MATRIX=blosum62 -TRANS=human40.cdi -LIST=45 -DCCALIGN=20 -THR_SCORB=pct
-THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext
-THR_MAX=100 -THR_MIN=0 -MAXLEN=2000000000
-HBAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
-USER=US10789129 @CGN 1 1 1549 @runat 23122005 152721 20318 -NCPU-6 -ICPU=3
-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Command line parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -MODEL=frame+
                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                               Score
      753
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seq length: 2000000000
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Match
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Ygapop 10.0,
Fgapop 6.0,
Delop 6.0,
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2:
3:
3:
5:
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753
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                                                                                                                                                                                                                                                                                                                                                                   /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
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/cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyright
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      926
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Ygapext
Fgapext
Delext
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  US-10-321-163-1

US-10-413-661-1

US-10-172-118-186

US-10-342-887-186

US-10-342-887-186

US-10-789-129-1

US-10-789-251-1

US-10-789-968-1
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7.0
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1452.183 Million cell updates/sec
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Sequence 1, Appli
Sequence 1, Appli
Sequence 186, App
Sequence 186, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
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ALIGNMENTS

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Query Match:
DB:
                                  Percent Similarity:
Best Local Similarity:
                                                                     Score:
                                                                                                     Alignment Scores:
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US-10-321-163-1
                                                                                     Pred. No.:
                                                                                                                                           US-10-321-163-1
                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Chandrasekher, Yasmin
APPLICANT: MCKETTAN, Patricia
APPLICANT: MCKETTAN, Patricia
TITLE OF INVENTION: METHOD FOR TREATING CERVICAL CANCER
FILE REFERENCE: 01-44US
CURRENT APPLICATION UNMBER: US/10/321,163
CURRENT FILING DATE: 2002-12-17
CURRENT FILING DATE: 2002-12-17
                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/10321163
Publication No. US20030161811A1
                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/341,783 PRIOR FILING DATE: 2001-12-17
                                                                                                                                                        ORGANISM: Homo sapiens FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)
                                                                                                                                                                                                                             TYPE: DNA
8.29e-93
753.00
100.00%
100.00%
100.00%
                                Length:
Matches:
Conservative:
Mismatches:
                   Indels:
926
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SOFTWARE: FastSEQ for Windows Version 3.
SEQ ID NO 1
LENGTH: 926
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)
US-10-413-661-1
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US-10-413-661-1
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                                                                                         Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                      Pred.
                                                                                                                                                                   Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/10413661
Publication No. US20030176657A1
GENERAL INFORMATION:
APPLICANT: Conklin, Darrell C.
APPLICANT: Haldeman, Betty A.
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10
                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 97-72
CURRENT APPLICATION NUMBER: US/10/413,661
CURRENT FILING DATE: 2003-04-15
PRIOR APPLICATION NUMBER: 09/199,586
PRIOR FILING DATE: 1998-11-25
PRIOR APPLICATION NUMBER: 06/066,597
PRIOR FILING DATE: 1997-11-26
NUMBER: OF SEQ ID NOS: 43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAGGAGACAGAA 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAACCTCAGGCAGCAGTTGTGAAGGCTTTGGGGGAACTAGACATTCTTCTGCAATGGATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CATTGTGGGGAAGGAATGAAGAAATACAGCCAGATTCTGAGTCACTTTGAAAAAGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCCAATTCCTTTCTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCCACATGACATGC
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                                                                                         8.29e-93
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                                                                         Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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GENERAL INFORMATION:

APPLICANT: Dai, Hongyue

APPLICANT: He, Yudong

APPLICANT: Hinsley, Peter

APPLICANT: Mao, Mao

APPLICANT: Mao, Mao

APPLICANT: Wan 't Veer, Laura

APPLICANT: Van 't Veer, Marc

APPLICANT: Van 't Vijver, Marc

APPLICANT: Dath Dagnosis and Prognosis of

FILE REFERENCE: 9301-175-99

CURRENT APPLICATION NUMBER: US/10/172,118

CURRENT FILING DATE: 2002-05-14

PRIOR APPLICATION NUMBER: 60/380,770

PRIOR FILING DATE: 2002-05-14
                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 2699

SEQ ID NO 186

LENGTH: 926

TYPE: DNA
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
DATABASE ECCESSION NUMBER: AF224266
DATABASE ENTRY DATE: 2001-06-18
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                                                    US-10-789-129-26 (1-144) x US-10-172-118-186
                                                                                               Query Match:
                                                                                                             Percent Similarity:
Best Local Similarity:
                                                                                                                                           Score:
                                                                                                                                                                    Alignment Scores:
                                                                                                                                                                                               US-10-172-118-186
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Publication No.
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CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
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753.00
100.00%
100.00%
100.00%
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Matches:
Conservative:
Mismatches:
Indels:
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APELICANT:
APPLICANT:
FILE NETHERENCE: 9301-188-999
CURRENT APPLICATION NUMBER: US/10/342,887
CURRENT PILING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: 60/298,918
PRIOR FILING DATE: 2001-06-18
PRIOR FILING DATE: 2002-05-14
PRIOR APPLICATION NUMBER: 60/380,710
PRIOR FILING DATE: 2002-05-14
PRIOR APPLICATION NUMBER: 10/172,118
PRIOR PILING DATE: 2002-05-14
PRIOR PILING DATE: 2002-06-14
VUMBER OF SEQ ID NOS: 2699
SEQ ID NO 186
LENGTH: 926
TYPE: DNA
ORGANISM: Homo sapiens
US-10-342-887-186
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Best Local Similarity:
Query Match:
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                                                                                      US-10-789-129-26 (1-144) x US-10-342-887-186 (1-926)
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APPLICANT:
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APPLICANT: He, Yudong
APPLICANT: Linsley, Pet
APPLICANT: Mao, Mao
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: Roberts, Christopher J.
: Van 't Veer, Laura Johanna
: Van de Vijver, Marc J.
: Bernards, Rene
                                                 CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer 20
ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
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Matches:
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Best Local Similarity:
Query Match:
DB:
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                                                                                                              US-10-789-129-26 (1-144) x US-10-789-129-1 (1-926)
                                                                                                                                                                                                                  Pred. No.:
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Publication No. US20040142428A1
GENERAL INFORMATION:
APPLICANT: Conklin, Darrell C.
APPLICANT: Haldeman, Betty A.
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10
FILE REFERENCE: 97-72
CURRENT APPLICATION NUMBER: US/10/789,129
CURRENT FILING DATE: 2004-02-27
                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 43
SOPTWARE: FastSEQ for Windows Version
SEQ ID NO 1
LENGTH: 926
                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/199,586
PRIOR FILING DATE: 1998-11-25
PRIOR APPLICATION NUMBER: 60/066,597
PRIOR FILING DATE: 1997-11-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/413,661
PRIOR FILING DATE: 2003-04-15
                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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             ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
                                                                     CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
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Gaps:
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APPLICANT: CONKLION.
APPLICANT: Haldeman, Betty A.
TITLE OF INVENTION: HAMMALIAN CYTOKINE-LIKE POI
FILE REFERENCE: 97-72
CURRENT APPLICATION NUMBER: US/10/789,251
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/413,661
PRIOR FILING DATE: 2003-04-15
PRIOR FILING DATE: 1998-11-25
PRIOR APPLICATION NUMBER: 09/199,586
PRIOR FILING DATE: 1998-11-25
PRIOR FILING DATE: 1998-11-26
PRIOR FILING DATE: 1997-11-26
NUMBER OF SEQ ID NOS: 43
SOPTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 926
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US-10-789-251-1
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; Publication No
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Publication No. US20040143097A1
GENERAL INFORMATION:
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ORCANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)
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                  GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC 320
                                  AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp 60
                                                                                GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
                                                                                                                                              TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGACATACGGGGCAGT 200
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APPLICANT: CONKIALOW.

APPLICANT: Haldeman, Betty A.

FILLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10

FILE REFERENCE: 97-72

CURRENT APPLICATION NUMBER: US/10/789,968

CURRENT FILING DATE: 2004-02-27

PRIOR APPLICATION NUMBER: US/10/413,661

PRIOR FILING DATE: 2003-04-15

PRIOR APPLICATION NUMBER: US/10/413,661

PRIOR FILING DATE: 2003-04-15

PRIOR APPLICATION NUMBER: 09/199,586

PRIOR APPLICATION NUMBER: 09/199,586

PRIOR FILING DATE: 1998-11-25

PRIOR APPLICATION NUMBER: 60/066,597

PRIOR FILING DATE: 1998-11-26

NUMBER OF SEQ ID NOS: 43

SOPTWARE: FASTSEQ for Windows Version 3.0

SEQ ID NO 1

LENGTH: 926

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

NAME/KEY. CIC
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US-10-789-968-1
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Publication No. US20040152878A1
GENERAL INFORMATION:
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                                           AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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Sequence 1, Application US/10748484

Publication No. US20040181040A1

GENERAL INFORMATION:
APPLICANT: Conklin, Darrell C.
APPLICANT: GENSMANN, Angelika
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10
FILE REFERENCE: 97-72
CURRENT APPLICATION NUMBER: US/10/748,484
CURRENT FILING DATE: 2003-12-29
PRIOR APPLICATION NUMBER: US/99/199,586
PRIOR FILING DATE: 1998-11-26
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 926
TYPE: DNA
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NAME/KEY: CDS
LOCATION: (45)...(572)
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                                           ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
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               AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
                                                                                    GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCCATTTGCTAAGACTCTATCTGGAC
                                                                                                AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
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US-10-994-116-1
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ORGANISM: Homo
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Sequence 1, Application US/10994116

Publication No. US20050136004A1

GENERAL INFORMATION:

APPLICANT: Xu, Wenfeng

APPLICANT: Kindsvogel, Wayne

APPLICANT: Chen, Zhi

APPLICANT: Chen, Zhi

APPLICANT: Lehner, Joyce M.

APPLICANT: Dillon, Stacey R.

APPLICANT: Lehner, Joyce M.

APPLICANT: Sivakumar, Pallavur V.

APPLICANT: Sivakumar, Pallavur V.

APPLICANT: Moore, Margaret D.

ITITLE OF INVENTION: ANTI-LL-20 ANTIBODIES AND BINDING PARTITILE OF INVENTION: INFLAMMATION

ITITLE OF INVENTION: NUMBER: US/10/994,116

CURRENT FILING DATE: 2004-11-19

PRIOR APPLICATION NUMBER: US 60/524,131

PRIOR APPLICATION NUMBER: US 60/525,857

PRIOR APPLICATION NUMBER: US 60/525,857
                                                                                                                                                                                                                                                                                                                            US-10-789-129-26 (1-144) x US-10-994-116-1 (1-926)
                                                                                                                ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
                                                                                                                                                                                                                                    CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                    GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
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Sequence 1, Application US/10994151

Publication No. US20050170468A1

GENERAL INFORMATION:

APPLICANT: Kindsvogel, Wayne
APPLICANT: Kindsvogel, Wayne
APPLICANT: Hughes, Steven D.
APPLICANT: Chandrasekher, Yasmin A.
APPLICANT: Chandrasekher, Yasmin A.
APPLICANT: Siadak, Anthony W.
APPLICANT: Moore, Margaret D.
TITLE OF INVENTION: ANTI-L1-20 RECEPTOR ANTIBODIES AND BINDING PARTNERS AND METHODS C.
TITLE OF INVENTION IN INFLAMMATION
FILE REFERENCE: 04-17

CURRENT APPLICATION NUMBER: US/10/994,151

CURRENT FILING DATE: 2004-11-19
PRIOR APPLICATION NUMBER: US 60/524,131

PRIOR APPLICATION NUMBER: US 60/555,857

PRIOR APPLICATION NUMBER: US 60/555,857

PRIOR APPLICATION NUMBER: US 60/555,857
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US-10-994-151-1
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; LOCATION: (45)...
US-10-994-151-1
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
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TYPE: DNA
ORGANISM: Homo
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                       GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCCATTTGCTAAGACTCTATCTGGAC
                                                                                                                                               ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
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TITLE OF INVENTION: AND EXPRESSION OF RECOMBINANT PROTEINS
FILE REFERENCE: 01-15
CURRENT APPLICATION NUMBER: US/10/122,882B
CURRENT FILING DATE: 2002-04-12
PRIOR APPLICATION NUMBER: 60/283,668
PRIOR FILING DATE: 2001-04-13
NUMBER: OF SEQ ID NOS: 25
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
LENGTH: 462
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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US-10-122-882B-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: POWNDER, TRACEY APPLICANT: CHAN, CHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          561
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81 AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
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                                                                           ArgValPheLy8AsnTyrGlnThrProAspHisTyrThrLeuArgLy8IleSerSerLeu
                                                                                                                                                     GACACCAAACCGGCGAACCGTTGCTGCCTGCCGCCCACCTGCTGCGCTCTGTATCTGGAC
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CURRENT FILING DATE: 2005-02-09
PRIOR APPLICATION NUMBER: 11/008,853
PRIOR FILING DATE: 2004-12-10
PRIOR APPLICATION NUMBER: US 60/529,412
PRIOR FILING DATE: 2003-12-12
PRIOR FILING DATE: 2003-12-12
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PastSEQ for Windows Version 4.0
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US-11-054-063-18
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LENGTH: 462
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APPLICANT: Pownder, Tracey
TITLE OF INVENTION: METHODS FOR ENHANCING
TITLE OF INVENTION: RECOMBINANT PROTEINS
FILE REFERENCE: 03-22C1
                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
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                                                                       AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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                            HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                      ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
                                                                                                                                                                  AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
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              CACTGTGGTGAGGAAGCAATGAAAAAATACAGCCAGATTCTGAGCCACTTCGAAAAAACTG
                                                                                                                                                      GACACCAAACCGGCGAACCGTTGCTGCCTGCTGCGCCACCTGCTGCTGTATCTGGAC
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TITLE OF INVENTION: AND EXPRESSION OF RECOMBINANT PROTBINS
FILE REFERENCE: 01-15
FILE REFERENCE: 01-15
CURRENT APPLICATION NUMBER: US/10/122,882B
CURRENT FILING DATE: 2002-04-12
PRIOR APPLICATION NUMBER: 60/283,668
PRIOR APPLICATION NUMBER: 60/283,668
PRIOR FILING DATE: 2001-04-13
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 513
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                 GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet
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 GAACCGCAGGCAGCAGTGGTGAAAGCTCTGGGTGAACTGGACATTCTGCTGCAGTGGATG
                                                                       CACTGTGGTGAGGAAGCAATGAAAAATACAGCCAGATTCTGAGCCACTTCGAAAAACTG
                                                                                              HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
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RESULT 14 US-10-467-431-1

GAGGAGACCGAA 510

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Sequence 1, Application US/10467431

| Sequence 1, Application VS/20040092445A1
| GENERAL INFORMATION:
| APPLICANT: Heuer, Josef
| APPLICANT: Heuer, Josef
| APPLICANT: Timothy, Noblitt
| TITLE OF INVENTION: USE OF LP82 TO TREAT HEMATOPOIETIC DISORDERS
| FILE REFERENCE: X.14753
| CURRENT APPLICATION NUMBER: US/10/467,431
| CURRENT FILING DATE: 2003-08-06
| PRIOR FILING DATE: 2001-02-28
| PRIOR APPLICATION NUMBER: 60/272,242
| PRIOR FILING DATE: 2001-11-19
| NUMBER OF SEQ ID NOS: 5
| SOFTWARE: Batentin version 3.1
| SEQ ID NO 1
| LENGTH: 867
| TYPE: DNA ORGANISM: Homo sapiens
| FEATURE: NAME/KEY: CDS | LOCATION: (272)..(799)
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| OTHER INFORMATION:
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NAME/KEY: sig_peptide
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OTHER INFORMATION:
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                     GluGluThrGlu 144
                                                   AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
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PRIOR FILING DATE: 2001-01-26
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.2
SEQ ID NO 1
LENGTH: 867
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publication No. US20040162237A1
GENERAL INFORMATION:
APPLICANT: FOX, Nyles W
TITLE OF INVENTION: USE OF LP82 TO TREAT BODY WEIGHT DISORDERS
FILE REFERENCE: X-14544
CURRENT APPLICATION NUMBER: US/10/466,695A
CURRENT FILING DATE: 2004-02-03
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (272)..(799)
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NAME/KEY: sig_peptide
LOCATION: (272)..(325)
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                                                                      121 GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet
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Job time : 823 secs

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